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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OM protein - protein search, using sw model
Published Applications AA:*

1: /cgn2=6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2=6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

3: /cgn2=6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

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254
1 TQAQLLRVGCVLGT
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         May 4, 2005, 18:14:26; Search time 134 Seconds (without alignments)
116.836 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Result No.	Score	Query Match Length	ength	BB	ID	Description
<b>,</b>	254	100.0	47	17	US-10-850-055-14	Sequence 14, App
N	72.5	28.5	52	φ	US-09-813-345-14	Sequence 14, A
ω	72.5	28.5	52	14	US-10-197-954-2	Sequence 2, A
4	72.5	28.5	52	16	US-10-474-635A-19	Sequence 19,
ហ	72.5	28.5	52		US-10-718-071-43	Sequence 43,
6	72.5	28.5	52		US-10-760-085-2	Sequence 2, A
7	72.5	28.5	185	15	US-10-364-889-6	Sequence 6, Appli
8	72.5	28.5	185		US-10-372-683-12	Sequence 12,
9	72.5	28.5	185		US-10-675-406A-7	Sequence 7, A
10	72.5	28.5	185		US-10-755-889-148	Sequence 148, App
11	69.5	27.4	50	9	US-09-813-345-15	Sequence 15, A
12	68.5	27.0	52	17	US-10-718-071-16	Sequence 16, Appl
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5	44	3	2	41	40	39	38	37	36	35	34	3	32	3	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14
53	53	53	•	53.5	53.5	53.5	54	54.5	54.5	54.5	55	55	55	56	56	56	56	56	56	56	56.5	57	58	58	58	8	8	58.5	59		63.5
20.9	.0		21.1	21.1	21.1	21.1	۲	21.5		:	:	21.7	:	22.0			22.0	,	2	22.0	'n	'n	'n	ν.	22.8	w	ω.	23.0	Ψ	24.6	25.0
136	117	98	329	303	303	135	538	751	320	126	669	322	220	947	107	107	107	107	107	89	575	117	1544	560	560	576	521	92	142	52	216
14	10	9	10	14	14	15	16	16	16	16	5	15	16	15	15	14	14	9	9	16	15	16	15	14	14	15	16	16	15	15	16
US-10-156-761-8699	-789-390-4	US-09-864-761-48541	-121	-171-311	-10-097-340	-264-049-	US-10-437-963-151317	Ψ	3	-10-437-963	-10-120-801	US-10-424-599-175411	-10-767-701-	194-11	-10-158-057	Ψ	US-10-125-540-356	US-09-764-853-470	US-09-764-870-356	US-10-437-963-136207	-10-42	-10-76	-10 - 36	-10-23	US-10-238-129-8	-10-184-648-	-10-437-963-12460	US-10-437-963-198935	US-10-307-817-14	US-10-360-101-74	US-10-437-963-198108
8699	e 41	485	φ,	e 87	ø	40				2020	108, A	æ	320	114	Sequence 335, App	301	e 356	470,	356,	136		51114,	60	8	Sequence 8, Appli	26	e 13	Sequence 198935,	14		Sequence 198108,

### ALIGNMENTS

WES-10-850-055-14 US-10-850-055-14 US-10-850-055-14 US-10-850-055-14 Sequence 14, Application US/10850055 Publication No. US20050009742A1 GENERAL INFORMATION: APPLICANT: Bertilsson, Goran APPLICANT: Brlandsson, Rikard APPLICANT: Heidrich, Jessica APPLICANT: Heidrich, Jessica APPLICANT: Heidrich, Johan APPLICANT: Heidrich, Johan APPLICANT: Lindquist, Per APPLICANT: Kortesman, Jarkko APPLICANT: Lindquist, Per APPLICANT: McGuire, Jacqueline APPLICANT: McGuire, Jacqueline APPLICANT: Nyberg, Karl APPLICANT: Nyberg, Karl APPLICANT: Ronnholm, Harriet APPLICANT: Ronnholm, Harriet APPLICANT: Wiketrom, Lillian APPLICANT: Wiketrom, Lillian APPLICANT: Wiketrom, Lillian APPLICANT: Zachrisson, Olof TITLE OF INVENTION COMPOUNDS FILE REFERENCE: 21882-517 CIP CURRENT APPLICATION NUMBER: US/10/850,055 CURRENT APPLICATION NUMBER: US/0/18,071 PRIOR APPLICATION NUMBER: US 00/127,912 PRIOR APPLICATION NUMBER: US 0/427,912 PRIOR FILING DATE: 2003-11-20 NUMBER OF SEQ ID NOS: 66 SOFTMARE: Patentin version 3.2 SEQ ID NO 14						<b></b> .	٠. ٠.	٠.	٠. ٠		٠		٠.	٠.	٠		٠.	٠.	٠		٠.	<b>S</b> 22
pplication US/108 ATTON: rtilsson, Goran rtilsson, Goran rtlandsson, Rikard risen, Jonas aegerstrand, Ande eldstrom, Kristin aggblad, Johan ansson, Katarina ortesman, Jarko indquist, Per undh, Hanna ccur, Jacquelin ercer, Alex yberg, Karl ssoinak, Amina atrone, Cesare onnholm, Harriet ikstrom, Lillian schrisson, Olof NTION: COMPOUNDS E: 2182-517 CUS G DATE: 2004-05- TION NUMBER: US 1 DATE: 2003-11-20 DATE: 2002-11-20 DATE: 2002-11-20 DATE: 2002-11-20 DATE: 2002-11-20 DD NOS: 66 entin version 3.2	I WE HER	PRIOR APP PRIOR FIL PRIOR APP	CURRENT A	TITLE OF	APPLICANT	APPLICANT	APPLICANT	APPLICANT	APPLICANT	APPLICANT	GENERAL IN	Sequence 1	RESULT 1 US-10-850-05									
	ING DATE: 2002-11-20 SEQ ID NOS: 66 PatentIn version 3 14 47	LICATION NUMBER: US 10/718, ING DATE: 2003-11-20 LICATION NUMBER: US 60/427,	PPLICATION NUMBER: US/10/85 FILING DATE: 2004-05-19			••		•••			•••		••	: Haggblad,	: Hellstrom,		••	••	g,	VEORMATION:	14, Application US/10850055	55-14
				INCREASING			•															
FOR INCREASING				NEUROGENESIS																		

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RESULT 3
US-10-197-954-2
US-10-197-954-2
US-20030119021A1
Publication No. US20030119021A1
                                                                                                                                                                                                                                              Query Match
Best Local Similarity
""" ches 15; Conserv
                                                                                                                                                                                                                                                                                                                                       SEQUENCE DESCRIPTION: SEQ ID NO: 14: US-09-813-345-14
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Sequence 14, Application US/09813345
Patent No. US20020068814A1
GENERAL INFORMATION:
APPLICANT: Smith, Derek D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        В
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Best Local Similarity
APPLICANT: K"ster, Hubert APPLICANT: Siddiqi, Suhaib
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 612/305-1220
TELEFAX: 612/305-1228
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 52 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: McCormack, Myra H
REGISTRATION NUMBER: 36,602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/813,345
FILING DATE: 20-War-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: peptid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 180.00020101 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MG-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Mueting, Raasch & Gebhardt, P.A.
STREET: 119 No. US20020068814A1th Fourth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: PEPTIDE ANTAGONISTS OF CGRP-RECEPTOR SUPERFAMILY AND METHODS OF USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                      15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       47;
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                                                                                                                                                                                                                        GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                                    GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: MN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOAQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 55401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Minneapolis
                                                                                                                                                                                                                                                               28.5%; Score 72.5; DB 9; Length 52; ilarity 38.5%; Pred. No. 0.067; Conservative 6; Mismatches 17; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 254; DB 17;
100.0%; Pred. No. 1.2e-25;
ative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                   Gaps
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APPLICANT:

Haegerstrand, Anders
Heidrich, Jessica
Hellstrom, Kristina
Hagblad, Johan
Jannson, Katarina
Kortesmaa, Jarkko

APPLICANT:

APPLICANT

APPLICANT: APPLICANT: APPLICANT:

> Erlandsson, Rikard Frisen, Jonas

APPLICANT:

APPLICANT: Bertilson, Goran

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APPLICANT: Isis Innovation Ltd
TITLE OF INVENTION: Peptides
FILE REFERENCE: 480821.00004
CURRENT APPLICATION NUMBER: US/10/474,635A
CURRENT FILING DATE: 2003-10-14
PRIOR APPLICATION NUMBER: GB 0109438.2
PRIOR FILING DATE: 2001-04-17
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn version 3.1
SEQ ID NO 19
LENGTH: 52
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                           밁
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                                                                                 RESULT 5
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                                                          US-10-718-071-43
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-474-635A-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/197,954
CURRENT FILING DATE: 2002-07-16
PRIOR APPLICATION NUMBER: 60/306,019
PRIOR FILING DATE: 2001-07-16
PRIOR APPLICATION NUMBER: 60/314,123
PRIOR APPLICATION NUMBER: 60/314,123
PRIOR FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: 60/363,433
PRIOR FILING DATE: 2002-03-11
NUMBER OF SEQ ID NOS: 149
NUMBER OF SEQ ID NOS: 149
Sequence 43, Application US/10718071 Publication No. US20050009847A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19, Application US/10474635A
Publication No. US20040176567A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 52
                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
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TITLE OF INVENTION: Capture Compounds, Collections Thereof
TITLE OF INVENTION: And Methods For Analyzing The Proteom
TITLE OF INVENTION: Compositions
FILE REFERENCE: 24743-2305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
                                                                                                                                           15
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                                                                                                                                                                                                                      15; Conservative
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                                                                                                                                         GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY 52
                                                                                                                                                                               GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
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                                                                                                                                                                                                                      6; Mismatches
                                                                                                                                                                                                                                         Score 72.5; DB 16; Pred. No. 0.067;
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                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                              52;
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Lindquist, Per

Hanna

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; SEQ ID NO 2
; LENGTH: 52
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-760-085-2
RESULT 7
US-10-364-889-6
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                                                                                                                                                                 Query Match
Best Local (
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                                                                                                                                                  Matches
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Publication No. US20050042771A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/760,085
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: 60/441,398
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 149
NOSTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/718,071
CURRENT FILING DATE: 2003-11-20
PRIOR APPLICATION NUMBER: US 60/427,912
PRIOR FILING DATE: 2002-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Capture Compounds, Collections Thereof
TITLE OF INVENTION: And Methods for Analyzing The Proteome
TITLE OF INVENTION: Compositions
FILE REFERENCE: 24743-2309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Hubert K"ster APPLICANT: Daniel Paul 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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SOFTWARE: PatentIn version 3.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                   Local Similarity
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                                                                                            9 GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
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T: Zachrisson, Olof
INVENTION: COMPOUNDS AND METHODS FOR INCREASING NEUROGENESIS
SRENCE: 21882-517 UTIL
                                                                         GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Suhaib Mahmood Siddiqi
Mattew Peter Grealish
Subramaniam Marappan
Chester Frederick Hassman
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Patrone, Cesare
Ronnholm, Harriet
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Nyberg, Karl
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38.5%;
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                                                                                                                                                                 Score 72.5; DB Pred. No. 0.067;
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                                                                                                                                                Mismatches
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                                                                                                                                                                                   DB 17; Length 52;
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Sequence 6, Application US/10364889

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.; ORGANISM: Homo sapien
US-10-372-683-12
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; ORGANISM: Homo sapiens
US-10-364-889-6
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SEQ ID NO 12
LENGTH: 185
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CURRENT FILING DATE: 2003-02-12
PRIOR APPLICATION NUMBER: 60/356,376
PRIOR FILING DATE: 2002-02-12
NUMBER OF SEQ ID NOS: 8
SOFTWARE: CuraSeqList version 0.1
SEQ ID NO 6
LENGTH: 185
                                                                                                                                                                                         Sequence 7, Application US/10675406A Publication No. US20040121375A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/675,406A
CURRENT FILING DATE: 2003-09-30
PRIOR APPLICATION NUMBER: US 60/415,194
                                                                             APPLICANT: Bayer Pharmaceuticals Corporation
APPLICANT: Eveleigh, Deepa
APPLICANT: Taylor, Ian
TITLE OF INVENTION: METHODS FOR PREDICTION AND
TITLE OF INVENTION: CANCER THERAPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/344,534 PRIOR FILING DATE: 2001-10-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/372,683
CURRENT FILING DATE: 2003-02-21
PRIOR APPLICATION NUMBER: US 10/271,690
PRIOR FILING DATE: 2002-10-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: GERRITSEN, MARY E. APPLICANT: PEALE JR., FRANKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Quinn, Kerry
TITLE OF INVENTION: Compositions and Methods for Treatment of Osteoarthritis
FILE REFERENCE: 21402-558
                                                               FILE REFERENCE: 5138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: METHODS FOR THE TREATMENT OF CARCINOMA FILE REFERENCE: P1928R1P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: PEALE JR., FR APPLICANT: WU, THOMAS D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Pahel, Gregory L. APPLICANT: Quinn, Kerry
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les 15; Conserv
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                                                                                                                                                                                                                                                                                                                                          109 GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY 146
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                                                                                                                                                                                                                                                                                                                                                                                                                            6; Mismatches
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                                                                                                     PROGNOSIS OF CANCER, AND MONITORING
                                                                                                                                                                                                                                                                                                                                                                                    47
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RESULT 11

US-09-813-345-15

; Sequence 15, Application U;

; Patent No. US20020068814A1

; GENERAL INFORMATION:
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LENGTH: 185;
TYPE: PRT;
ORGANISM: Homo sapiens
US-10-675-406A-7
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APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: POLYMUCLEOTIDES AND POLYPEPTIDES ASSOCIATED WITH THE NF-KB
TITLE OF INVENTION: PARHWAY
FILE REFERENCE: D0284 NP
CURRENT FILING DATE: 2004-01-13
PRIOR APPLICATION NUMBER: U.S. 60/440,068
PRIOR APPLICATION NUMBER: U.S. 60/440,068
PRIOR APPLICATION NUMBER: U.S. 60/449,757
PRIOR APPLICATION NUMBER: U.S. 60/469,757
PRIOR APPLICATION NUMBER: U.S. 60/469,757
PRIOR FILING DATE: 2003-05-12
NUMBER OF SEQ ID NOS: 823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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SEQ ID NO 148
LENGTH: 185
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NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
les 15; Conserv
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               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEB: Mueting, Raasch & Gebhardt, P.A.
STREET: 119 No. US20020068814A1th Fourth Street
                                                                                                                                                                                                                                                             TITLE OF INVENTION: PEPTIDE ANTAGONISTS OF CGRP-RECEPTOR
                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                     APPLICANT: Smith, Derek D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      109 GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             109 GCRFGTCTVOKLAHQIYOFT-DKDKDNVAPRSKISPOGY 146
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                                  STATE: MN
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                                                                                                 ZIP: 5540:
                                                                                                               COUNTRY: USA
                                                                                                                                                CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                               Application US/09813345
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5. US20040171823A1
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Abel, Peter W.
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PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28.5%;
                                                                                                                                                                                                                                                SUPERFAMILY
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Pred. No. 0.27;
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                                                                                                                                                                                                                                              AND METHODS OF USE
Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17;
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; LENGTH: 52
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-718-071-16
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Best Local S
Matches 14
        Matches
                                                                                                                              SOFTWARE: PatentIn version 3.2 SEQ ID NO 16
                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 publication No
                                                                                                                                                                           APPLICANT: Zachrisson, Olof
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR INCREASING NEUROGENESIS
FILE REFERENCE: 21882-517 UTIL
CURRENT APPLICATION NUMBER: US/10/718,071
CURRENT FILING DATE: 2003-11-20
PRIOR APPLICATION NUMBER: US 60/427,912
PRIOR FILING DATE: 2002-11-20
PRIOR FILING DATE: 2002-11-20
                                                                                                                                                              NUMBER OF SEQ ID NOS: 71
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TELEPAX: 612/305-1228
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/813,345
FILING DATE: 20-Mar-2001
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13 GCRFGTCTMQKLAHQIYQFT-DKDKDGMAPRNKISPQGY 50
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                     Similarity
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Erlandsson, Rikard
Frisen, Jonas
Haegerstrand, Anders
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Mercer, Alex
Nyberg, Karl
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TYPE: amino acid
STRANDEDNESS: single
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REFERENCE/DOCKET NUMBER: 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/10718071
No. US20050009847A1
                                                                                                                                                                                                                                                                                                          Ronnholm, Harriet
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Patrone, Cesare
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Jannson, Katarina
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     Conservative
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Lindquist, Per
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hellstrom, Kristina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      feidrich, Jessica
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                27.0%;
38.5%;
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                  Score 68.5;
Pred. No. 0
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   Mismatches
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                                DB 17;
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 Indels
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US-10-264-237-1622
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SEQ ID NO 198108
LENGTH: 216
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Best Local :
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                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                          Sequence 198108, Application US/10437963 Publication No. US20040123343A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                               TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
                                  NUMBER OF SEC ID NOS: 204966
                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                  APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
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OTHER INFORMATION: Xaa equals any of the twenty naturally occurring L-amino
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NAME/KEY: MISC_FEATURE
LOCATION: (67)
OTHER_INFORMATION: Xaa
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Pred. No. 1.8;
5; Mismatches
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Search completed: May
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SEQ ID NO 74
LENGTH: 52
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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APPLICANT: Moll, Gert N.
APPLICANT: Leenhouts, Cornelis J.
TITLE OF INVENTION: Export and modification of (poly) peptide in the lantibiotic way
FILE.REFERENCE: 2183-5673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 74, Application US/10360101
Publication No. US20040009550A1
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                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 309
                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/360,101
CURRENT FILING DATE: 2003-02-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Oryza sativa
                                                                                                                                                                                                      OTHER INFORMATION: Al3, S16-sequence of Adrenomedullin Hypotensive
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                                                                                                                                           Similarity
                                                            GSRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY
                                                                                          GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
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              4, 2005,
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35.9%;
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Pred. No. 4.
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Pred. No. 1
                                                                                                                            Mismatches
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Post-processing: Minimum Match 0%
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1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
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254
1 TQAQLLRVGCVLG
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geneseqp2003bs:*
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112.208 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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Aab75124 UspA(1-84	Abj18670 Universal			UspA (1-	Adf55556 Active hu	Aab75113 Linker pe	Abj18665 Universal		Aab75111 Glycine e			Adp18429 Neurogene	Adm98309 Mature hu	Adn03278 Exemplary	Human	Human	Human	Abp55104 Human adr	Human	Aab75110 Human adr	Aab91759 Adrenomed		Ado61520 Human int	Ado61523 Human int	Description	

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70.5	71.5	71.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	72.5	
27.8	28.1	28.1	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	28.5	
188	184	184	206	186	185	185	185	185	185	185	185	185	185	185	185	185	185	170	
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AAB60345	ADO60040	ABB57209	ABJ18668	ADD18582	ADS19162	ADR87608	ADR14147	AD036937	ADP12587	ADP19855	ADM98308	ADN10849	ADF55557	ADA27595	ABP72347	AAB49697	AAB60344	AAB75114	PHOTOCAL
Aab60345	Ado60040	Abb57209	Abj18668	Add18582	Ads19162	Adr87608	Adr14,147	Ado36937	Adp12587	Adp19855	Adm98308	Adn10849	Adf55557	Ada27595	Abp72347	Aab49697	Aab60344	Aab75114	1,004
Porcine a	) CRH signa	Mouse isc	Universal	Human	! Human adr	Human	Human	' Human pro	Protein e	Human	Human pre	Human adr	Human adr	Human	Adrenomed	' Human adr	Human adr	Thioredox	01111

ALIGNMENTS

RESULT 1 AD061523

Human intermedin mature proetin SEQ ID NO:5. ovarian follicle human; intermedin; hypotensive; gene therapy; hypertension; cardioprotective; diet; prolactin release; growth hormone release; 26-AUG-2004 (first entry) ADO61523; ADO61523 standard; protein; 47 26-NOV-2003; 2003WO-US037968 10-JUN-2004. WO2004048547-A2 Homo sapiens. survival; oedema A

(STRD ) UNIV LELAND STANFORD JUNIOR.

26-NOV-2002; 2002US-0429327P

Hgu SYT;

WPI; 2004-441176/41.

New composition comprises an intermedin peptide, useful for treating hypertension, as cardioprotective agent, as a diet aid, for the release of prolactin, or for growth hormone release from the pituitary.

Disclosure; SEQ ID NO 5; 68pp; English.

The invention relates to a novel composition comprising an intermedin peptide, which comprises at least 18 contiguous amino acids in a sequence comprising 148 amino acids (ADOSIS20). A composition of the invention has hypotensive activity, and may have a use in gene therapy. The composition comprising the intermedin peptide is useful for identifying homologous or related genes, for production of the encoded peptide, in producing compositions that modulate the expression or function of its encoded protein, for gene therapy, mapping functional regions of the protein, or in studying associated physiological pathways. The intermedin peptide is useful for treating hypertension, as a cardioprotective agent, as a diet aid, for the release of protactin, in growth hormone release from the pituitary, or for ovarian follicle survival and growth. They are also

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TQAQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 147

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RESULT 2
ADO61520
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   Matches
                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                  peptide, which comprises at least 18 contiguous amino acids in a sequence comprising 148 amino acids (ADO61520). A composition of the invention has hypotensive activity, and may have a use in gene therapy. The composition comprising the intermedin peptide is useful for identifying homologous or related genes, for production of the encoded peptide, in producing compositions that modulate the expression or function of its encoded protein, for gene therapy, mapping functional regions of the protein, or in studying associated physiological pathways. The intermedin peptide is useful for treating hypertension, as a cardioprotective agent, as a diet aid, for the release of prolactin, in growth hormone release from the pituitary, or for ovarian follicle survival and growth. They are also useful in the reduction of oedema. The present sequence represents the
                                                                 Sequence 148 AA;
                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 2; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                 New composition comprises an intermedin peptide, useful for treating hypertension, as cardioprotective agent, as a diet aid, for the release of prolactin, or for growth hormone release from the pituitary.
                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-441176/41.
N-PSDB; ADO61519.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hsu SYT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; intermedin; hypotensive; gene therapy; hypertension; cardioprotective; diet; prolactin release; growth hormone release; ovarian follicle survival; oedema.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-NOV-2002; 2002US-0429327P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004048547-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human intermedin SEQ ID NO:2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADO61520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADO61520 standard; protein; 148 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful in the reduction of oedema. The present sequence represents the human intermedin mature protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-NOV-2003; 2003WO-US037968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-AUG-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (STRD ) UNIV LELAND STANFORD JUNIOR
                  Local
                                                                                             intermedin protein of the invention.
 l Similarity 47; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 TOAQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      47; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TQAQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   47 AA;
   Conservative
                                                                                                                                                                                                                                                                                                   relates to a novel composition comprising an intermedin
             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
0
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Score 254; DB 8;
Pred. No. 3.9e-27;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 254; DB 8;
Pred. No. 1.1e-27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ٥,
                              Length 148;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
Indels
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0
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Gaps /
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0
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TQAQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY

RESULT 4 AAB91759

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RESULT 3
AAB91765
ID AAB9
XX AAB9
XX AAB9
XX AAB9
XX AAB9
XX AAB9
XX Adre
XX Prot
KW bloo
KW bloo
KW hydr
OS Homo
OS Synt
XX 17-M
PF 17-M
PF 17-M
PR 11-S
PF 18-G
PF 15-G
XX 1
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                                                      S
                                                                                                                                                                                                                                                                                      The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a CC reactive group (II) (e.g. succinimidyl and maleimido groups) attached to CC a less therapeutically active amino acid region (IV), which covalently CC bonds with amino/hydroxyl/thiol groups on blood components to form a CC peptidase stabilised therapeutic peptide composed of 3-50 amino acids. CC (I) are useful for modifying therapeutic peptides e.g. hormones, growth CC factors and neurotransmitters, to protect them from peptidase activity in CC vivo for the treatment of various disorders. Endogenous therapeutic CC peptides are not suitable as drug candidates as they require frequent CC administration due to rapid degradation by peptidases in the body. CC Modifying and attaching therapeutic peptides to albumin prevents or CC reduces the action of peptidases to increase length of activity (half CC life) and specificity as bonding to large molecules decreases CC intracellular uptake and interference with physiological processes. CC AAB90829 to AAB92441 represent peptides which can be used in the CC exemplification of the present invention
                                                                                                                         Matches
                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-MAY-1999;
10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity.
                                                                                                                                                                                                                                            Sequence 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 501;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-112059/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bridon DP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200069900-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Adrenomedullin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB91765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CONJ-) CONJUCHEM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-MAY-2000; 2000WO-US013576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino
   14
                                                            9
                                                                                                                      14;
                                                                                                                                                   Similarity
GCRFGTCTVQKLAHQIYQFTDKDGVAPRSKISKISPQGY 52
                                                         GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      thiol;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ezrin AM,
                                                                                                                                                                                                                                            AΑ;
                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0134406P.
99US-0153406P.
99US-0159783P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hormone; growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide;
                                                                                                                                                29.9%;
35.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            733pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Milner PG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AM)
                                                                                                                   9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID
                                                                                                                                                Score 76; DB 4; Length 52; Pred. No. 0.0091;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO:941.
                                                                                                                   19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thibaudeau
                                                                                                                   Indels
                                                                                                                   0
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                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                 The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a CC reactive group (II) (e.g. succinimidy] and maleimido groups) attached to call the group (II) and a stacked to compose the repeutically active amino acid region (IV), which covalently composed to stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in composed of 3-50 amino acids. (C) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in composed of 3-50 amino acids. (C) expides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. (C) Modifying and attaching therapeutic peptides to albumin prevents or composed the action of peptidases to increase length of activity (half if) and specificity as bonding to large molecules decreases contracellular uptake and interference with physiological processes. (C) AAB90829 to AAB92441 represent peptides which can be used in the contractive contracellular uptake and interference with physiological processes.
                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 15
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10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protection; endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
               31-JUL-2001
                                                 AAB75110;
                                                                                 AAB75110 standard; protein;
                                                                                                                                                                                                                                                                                                       Sequence 52 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bridon DP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 498; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-112059/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-MAY-2000; 2000WO-US013576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-NOV-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200069900-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Adrenomedullin peptide (AM) SEQ ID NO:935.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB91759 standard; peptide; 52
                                                                                                                                                                      15
                                                                                                                                                                                                                                     1 Similarity
15; Conserv
                                                                                                                                                                                                    9
                                                                                                                                                                    GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY 52
                                                                                                                                                                                                    GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ezrin AM,
                                                                                                                                                                                                                                   28.5%;
ilarity 38.5%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
              (first entry)
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99US-0153406P.
99US-0159783P.
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                                                                                 52
                                                                                                                                                                                                                                     6
                                                                                                                                                                                                                                   Score 72.5; DB
Pred. No. 0.028;
6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Holmes DL,
                                                                                                                                                                                                                                                                  DB 4;
                                                                                                                                                                                                                                     17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thibaudeau
                                                                                                                                                                                                                                                                     Length 52;
                                                                                                                                                                                                                                     Indels
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ARESULT 6
AAEOJAE
ID AAEO
XX
AC AAEO
XX
AC AAEO
XX
DT 29-N
XX
DT Uuma
XX
Huma
XX
CGRP
XX
US62
XX
OS Homo
XX
OS Homo
XX
OS JOA
XX
PF 30-A
XX
PF 30-A
XX
PR 30-A
XX
PA (UYC
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Best Local Similarity
Matches 15; Conserv
                                                                                                                                                       Human; vasoactive peptide; calcitonin gene related peptide; CGRP-
CGRP-receptor identification; adrenomedullin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 52 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dissolving the sediment and extracting adrenomedullin precursor. The method can be used for the production of adrenomedullin precursor for pharmaceutical and diagnostic applications. AAH19806 to AAH19866 and AAB75110 to AAB75124 represent sequences which are used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Producing adrenomedullin useful application comprises producing recombinant host.
                          30-APR-1998;
                                                    30-APR-1998;
                                                                              31-JUL-2001
                                                                                                                               Homo sapiens
                                                                                                                                                                                              Human adrenomedullin peptide
                                                                                                                                                                                                                          29-NOV-2001
                                                                                                                                                                                                                                                                           AAE09818 standard; peptide; 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a method (M1) for producing adrenomedullin precursor. The method comprises: (a) producing the fused protein using a recombinant host cell; (b) restricted digestion of the fused protein by a protease followed by collection of sediment; and (c)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Adrenomedullin; glycine extended adrenomedullin; AM; AM-gly; adrenomedullin precursor; fusion protein; pharmaceutical; diagnostic.
                                                                                                       US6268474-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-282044/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Takimoto A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human adrenomedullin (AM) protein
 (UYCR-) UNIV CREIGHTON.
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                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99JP-00294147
                          98US-00070504
                                                    98US-00070504.
                                                                                                                                                                                                                                                                                                                                                                                                           28.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             75pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nakayama T,
                                                                                                                                                                                                                                                                                                                                                                                             Score 72.5; DB Pred. No. 0.028; 6; Mismatches
                                                                                                                                                                                                #
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for pharmaceutical and diagnostic fused adrenomedullin precursor using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mitsushima K;
                                                                                                                                                                                                                                                                                                                                                                                                                        DB 4;
                                                                                                                                                                                                                                                                                                                                                                                               17;
                                                                                                                                                                                                                                                                                                                                                                       47
                                                                                                                                                                                                                                                                                                                                             52
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RESULT 7
ABP55104
ID ABP55104
ID ABP55104
ID ABP55104
ID O7-E
XX ABP5
XX ADR6
XX Adr6
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Best Local S
Matches 15
                         The present sequence is that of human adrenomedullin, a potent hypotensive and vasodilator. The invention provides modified calciton and related peptides, such as adrenomedullin, that have at least 70% identity to the native form but are modified such that the tendency of the peptide to aggregate is reduced. Preferred regions for modification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to antagonists of the vasoactive peptide calcitonin gene related peptide (GGRP) and other members of the GGRP superfamily. The invention also relates to amino the terminal modifications of peptides to improve their ability to bind to a member of the CGRP receptor super-family. CGRP antagonists are used for inhibiting CGRP activity which can be used in vitro e.g. in assays to identify and/or isolate CGRP receptors or with intect cells either in vitro or in vivo to inhibit the effect of CGRP binding to its receptor. The present sequence
                                                                                                                                                                                                   Example;
                                                                                                                                                                                                                                                                                                                                                                                                 Zurdo J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-APR-2001; 2001GB-00009438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200283734-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Adrenomedullin; human; pro hypotensive; vasodilator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human adrenomedullin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vasoactive peptides useful for inhibiting calcitonin gene related peptide receptor activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-APR-2002; 2002WO-GB001778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABP55104 standard; peptide; 52 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 5; Col 25-26; 24pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Smith
                                                                                                                                                                                                                                              modified human calcitonin peptide having reduced aggregation, useful the treatment of Paget's disease, hypercalcemia and/or osteoporosis.
                                                                                                                                                                                                                                                                                                                                          2003-046916/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15
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15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                             Page 21; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   adrenomedullin peptide
                                                                                                                                                                                                                                                                                                                                                                                                 Dobson CM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       INNOVATION LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ŝ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers 16. .21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein engineering;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6
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Pred. No. 0.028;
6; Mismatches 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           solubility;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           aggregation;
                               tendency of modification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ۲,
                                                                                                                calcitonin
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15

GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY

52

v 15;

Matches Query Match Best Local

Similarity

28.5%; ilarity 38.5%; Conservative

Score 72.5; DB Pred. No. 0.028; 6; Mismatches

17;

Indels

1;

Gaps

DB 7;

Length 52;

Sequence 52

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RESULT 8
ADC25152
ID ADC22
XX ADC2
XX ADC2
XX ADC2
XX Huma
XX Cytc
XX Iive
XX Homc
XX The
CC Gur
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 15
                                                                                      compositions for treating cancer e.g. stomach cancer, colon cancer, pulmonary cancer, ovarian cancer, liver cancer or pancreatic cancer. current sequence represents an angiogenesis inhibiting peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                            The invention relates to peptides comprising an amino acid sequence derived from a fully defined 52 amino acid sequence (S1) given in the specification. Peptides may be created by deletion of some amino aci from the N-terminal of S1. The peptides are applicable in drug
                                                                                                                                                                                                                                                                                                                                                                                     Peptides with effect on inhibiting angiogenesis in cancer cells and inhibiting proliferation of cancer cells, and encoded polynucleotides, applicable in drug compositions for treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           include those for which the peptide is polymorphic amongst different species, which increase the propensity of the peptide to form local interactions of the alpha-helical type, or which reduce the number of hydrophobic residues or increase the net charge of the peptide. When aggregation is reduced or prevented, lower doses of the drug can be us Side-effects and undesired responses are minimised by retaining high sequence identity to the human peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-767505/72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-MAR-2002; 2002JP-00075575.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-MAR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HOKK-) HOKKAIDO TECHNOLOGY LICENSING OFFICE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003078460-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human angiogenesis inhibiting peptide #SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC25152;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC25152 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                          SEQ ID NO 1; 41pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy; cancer; stomach;
reatic; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2003WO-JP003344.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide; 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 72.5; D. Pred. No. 0.02 6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72.5;
No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     colon;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pulmonary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     52
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                                                                                               of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                              acids
                                                                                                                        The
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RESULT 10
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#XXX
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                                                                                                                                                                                                                                                                                                                                  The invention relates to a method of screening for schizophrenia in a population comprises determining the magnitude of expression, in members of the population, of at least one gene selected from the gene encoding decidual protein induced by progesterone (DEPP), the gene encoding adrenomedullin and the gene encoding cold shock domain protein A (csdA) in a sample, and comparing the magnitude of expression to a baseline of expression of the gene, where increased gene expression indicates the presence of schizophrenia. An antisense molecule, siRNA, ribozyme or nucleic acid molecule promoting triple helix formation that specifically inhibit the expression of DEPP, csdA or adrenomedullin genes, is useful for the manufacture of a medicament for the treatment of schizophrenia. An antibody that specifically binds an epitope of DEPP, csdA or adrenomedullin is also useful for the manufacture of a medicament for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Screening for schizophrenia in a population utilizing genes encoding the decidual protein induced by progesterone, adrenomedullin and/or cold shock domain protein A, useful in diagnosing, preventing and/or treating
              ADM94034
                                              ADM94034 standard; peptide; 52 AA
                                                                                                                                                                                                                                                                      Sequence 52 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 1; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-767532/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Buxton FP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-MAR-2003; 2003WO-EP002875
                                                                                                                                                                                                                                                                                                    adrenomedullin protein.
                                                                                                                                                                                                                                                                                                                       treatment of schizophrenia. This sequence represents the mature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAR-2002; 2002US-0366001P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003078658-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sirna;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neuroleptic; antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mature adrenomedullin protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NOVS ) NOVARTIS AG. (NOVS ) NOVARTIS PHARMA GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-SEP-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DEPP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG91993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG91993 standard; protein; 52 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              leptic; antisense therapy; gene therapy; adrenomedullin agonist; ophrenia; gene expression; decidual protein induced by progesterone; adrenomedullin; cold shock domain protein A; csdA; antisense;
                                                                                                                                  15
                                                                                                                                                                                                    15;
                                                                                                                                                                  v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ribozyme; triple helix formation
                                                                                                                                                                                                                     Similarity
                                                                                                                                                              GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                              GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Carpenter WT,
                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                  28.5%;
38.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberts RC,
                                                                                                                                                                                                    6;
                                                                                                                                                                                                                  Score 72.5; DB 7; Pred. No. 0.028;
                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tamminga CA;
                                                                                                                                                                                                  17;
                                                                                                                                                                                                                                Length 52;
                                                                                                                                                                                                    Indels.
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RESULT 11
ADN03278
ID ADN03
XX
AC ADN03
XX
DT 17-JU
XX
DE Exemp
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                                                                                                                                                                                                                                                                                                                                                                  CC (CRSPs) that are expressed in the central nervous system and act on the CC calcitonin receptor to promote the production of cAMP within a cell. CC Specifically, it refers to CRSPs that can stimulate concentration CC dependent sodium ion uptake and furthermore act to inhibit calcium ion CC dependent sodium ion uptake and furthermore act to inhibit calcium ion CC dependent sodium ion uptake and furthermore act to inhibit calcium ion CC derived from the CRSPs that have additions, deletions and/ or CC derived from the CRSPs that have additions, deletions and/ or CC desire thions of one or more amino acids, yet that retain similar CC activity and are useful for developing drug compositions and CC pharmaceutically acceptable carriers to treat and/ or prevent skeletal CC disorders, cancer, hypertension and restenosis. Furthermore, CRSPs can CC also be used as analgesics, appetite suppressants and diuretics such that CC exhibit vasotropic, cytostatic, diuretic, osteopathic, anorectic and CC hypotensive activities. This peptide sequence is a human AM peptide that CC has cAMP activity, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                     Query Match
                                   17-JUN-2004
                                                                                                                                                                                                                                                                                                                                          Sequence 52 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Calcitonin receptor stimulating peptides promoting cAMP production cells for treatment of cancer, skeletal disorders and hypertension appetite suppressants and analgesics.
                                                                                                       ADN03278 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 2; 62pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   calcitonin receptor stimulating peptide; CRSP; cAMP activity; skeletal disorder; cancer; hypertension; restenosis; analgesic; appetite suppressant; diuretic; vasotropic; cytostatic; diuretic; osteopathic; anorectic; hypotensive; human; AM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to novel calcitonin receptor stimulating peptides (CRSPs) that are expressed in the central nervous system and act on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-043113/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Minamino N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-JUN-2002; 2002JP-00162797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-MAY-2003; 2003WO-JP006641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003102180-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disulfide-bond Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human AM peptide, a CRSP homologue.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NISC-) JAPAN SCI & TECHNOLOGY CORP. (NINA-) JAPAN NAT CARDIOVASCULAR.
                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                               15
                                                                                                                                                                                                                                9
                                                                                                                                                                                                                                                                   15;
                                                                                                                                                                                                                              GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                                                                                           GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY
                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Katafuchi T;
                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
16. .21
52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "C-terminal amide"
                                                                                                      peptide;
                                                                                                                                                                                                                                                                                    28.5%;
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و
                                                                                                                                                                                                                                                                                    Score 72.5; DB Pred. No. 0.028;
                                                                                                       B
                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                   DB 8; Length
                                                                                                                                                                                                                                                                   17;
                                                                                                                                                                                             52
                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                      52
                                                                                                                                                                                                                                                                   1;
                                                                                                                                                                                                                                                                   Gaps
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and as
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Exemplary peptide ligand for proteome analysis

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SAXAXE
                                                                                                                RESULT 12
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                                                                                                                                                                                                                                                                                                                                            The invention relates to a collection of capture compounds capable of CC binding to biomolecules to form complexes that are stable under mass comperisors of compounds of formula (I)-(III) given in the specification. Also CC sets of compounds of formula (I)-(III) given in the specification. Also CC included are analysis of biomolecules (by contacting a composition comprising a biomolecule with the above collection and identifying or CC detecting bound biomolecules), separating protein conformers (by CC contacting a composition comprising a biomolecule with the above collection, separating the members of the collection and identifying or CC contacting a composition comprising a biomolecule with the above collection and separating each count groteins), reducing diversity of a complex mixture of biomolecules (by contacting the mixture with the above collection and separating each contacting the complexes of capture compounds with biomolecules (by sorting cells from a single subject into sets according to a phenotype, contacting comparing the patterns of biomolecule binding from each set). The CC contacting of capture compounds is useful for the analysis of biomolecules, especially proteins (e.g. analysis of a proteome), using contacting of flight (MALDI-TOP) mass spectrometry. The present sequence is an exemplary peptide ligand which may be incorporated into a capture
                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                   Query Match
                 15-JUL-2004
                                                 ADM98309;
                                                                                ADM98309 standard;
                                                                                                                                                                                                                                                                                                      Sequence 52
                                                                                                                                                                                                                                                                                                                                   exemplary peptide compound of the in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Collection of capture compounds capable of binding to biomolecules to form complexes that are stable under mass spectrometry conditions, useful for analysis of biomolecules, especially proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-JUL-2001; 2001US-0306019P.
21-AUG-2001; 2001US-0314123P.
11-MAR-2002; 2002US-0363433P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                matrix assisted laser desorption ionisation-time of flight; MALDI-TOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; SEQ ID NO 2; 165pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-059185/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Koster H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KOST/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-JUL-2002; 2002US-00197954.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2003119021-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein separation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide ligand; proteome; capture compound; mass spectrometry;
                                                                                                                                                                                                                                                      Local
                                                                                                                                                                  15
                                                                                                                                                                                                                                   15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ) KOSTER H.
) SIDDIQI S.
) LITTLE D P.
                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                  GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                 GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Siddiqi S,
                                                                                                                                                                                                                                                                                                      A,
                                                                                                                                                                                                                                   Conservative
              (first entry)
                                                                                                                                                                                                                                                                                                                                     invention.
                                                                                protein;
                                                                                                                                                                                                                                                   28.5%;
38.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Little DP;
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                                                                                                                                                                                                                                   <u>٠</u>.
                                                                                                                                                                                                                                                   Score 72.5;
Pred. No. 0
                                                                                B
                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                     0.028;
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                                                                                                                                                                                                                                                                 8;
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RESULT 13
ADP18429
ID ADP18
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to a novel peptide AM(11-22) useful for treating shock, or in a pharmaceutical composition for inducing vasoconstriction. Specifically, AM(11-22) is a short peptide derived from human adrenomedullin (AM), which is a vasoregulatory compound that influences blood pressure. The present invention describes screening assays to identify compounds including antibodies, small molecule inhibitors or peptides that modulate AM(11-22)-mediated vasoconstriction and as such represent novel vasodilators or vasoconstrictors. Accordingly, AM(11-22) can be used therapeutically in a pharmaceutical composition to inhibit blood flow following traumatic or surgical injury, as well as for vasodilatory, septic or haemorrhagic shock, and thus exhibits vasotropic, hypotensive, immunosuppressive and antibacterial activities. This polypeptide sequence is the mature human adrenomedullin protein of the
          haemostatic; hypertensive; muscular-Gen; ophthalmological antiinflammatory; analgesic; antidiabetic; neurogenesis m
                                           neurodegenerative; ischaemic; learning and memory disorder; neurological trauma; nootropic; neuroprotective; CNS-Gen; cerebroprotective; vasotropic; anticonvulsant; antiparkinsonian;
                                                                                                                             Neurogenesis modulation-related protein SeqID43.
                                                                                                                                                              26-AUG-2004
                                                                                                                                                                                               ADP18429;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 52
                                                                                             neurogenesis modulation; neural tissue;
                                                                                                                                                                                                                            ADP18429 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-OCT-2002; 2002US-0416291P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-OCT-2003; 2003WO-US031400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 3; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New AM(11-22) peptides, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Saavedra JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004032708-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                haemorrhagic shock; vasotropic; hypotensive; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AM(11-22); vasoconstriction; human; adrenomedullin; AM; vasoregulatory; blood pressure; vasodilator; vasodilatory shock; septic shock;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mature human adrenomedullin protein SeqID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treating septic shock, vasodilatory reducing blood pressure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2004-340778/31.
                                                                                                                                                                                                                                                                                                               15
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                                                                                                                                                                                                                                                                                                                                                9
                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                              GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY
                                                                                                                                                                                                                                                                                                                                              GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                               A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Martinez
                                                                                                                                                             (first entry
                                                                                                                                                                                                                                                                                                                                                                                             28.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ņ,
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                                                                                                                                                                                                                                52
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                                                                                                                                                                                                                                                                                                                                                                             Score 72.5; DB 8;
Pred. No. 0.028;
6; Mismatches 17
                                                                                                                                                                                                                                A
A
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                                                                                             central nervous system disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for inducing vasoconstriction, nock or hemorrhagic shock, or
                                                                                                                                                                                                                                                                                                                                                                                                              Length 52;
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'neural tissue G-protein coupled receptor activator; neurogenesis inducer; intracellular neural cAMP enhancer; intracellular neural cAMP enhancer; intracellular neural camp enhancer; intracellular neural camp enhancer; Parkinson's disease; Parkinson's disease; Huntington's disease; Alzheimer's disease; multiple sclerosis; amyotrophic lateral sclerosis; Shy-Drager syndrome; progressive supranuclear palay; Lewy body disease; spinal ischaemia; ischaemic stroke; cerebral infarction; spinal cord injury; cancer-related brain; spinal cord injury; multi-infarct dementia; geriatric dementia; cAMP level; embryonic tissue; human.
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WO2004045592-A2

20-NOV-2003; 2003WO-IB005311.

20-NOV-2002; 2002US-0427912P

SEERE (JANS/ (KORT/ (LIND/ (HEID/ (FRIS, (ERLA/) NEURONOVA AB. BERTILSSON G. ERLANDSSON R. HAEGGBLAD J. JANSSON K. KORTESMAA J. LINDQUIST P. HEIDRICH J. HELLSTROEM K. HAEGESTRAND A. FRISEN J.

(NJBE/ (OSSO/ (PATR/ (TUND/ MERC OSSOINAK A.
PATRONE C.
ROENNHOLM H.
ZACHRISSON O.
WIKSTROEM L. MERCER A. NJBERG K. LUNDH H. MCGUIRE J.

Hellstroem Lundh H, M Bertilsson Roennholm n G, Erlandsson R, m K, Haeggblad J, Mcguire J, Mercer Zachrisson O, , Frisen J, Ha,
Jansson K, Ko,
r A, Njberg K,
Wikstroem L; Haegestrand A, Heidrich J Kortesmaa J, Lindquist P; K, Ossoinak A, Patrone C; Heidrich J

(WIKS/)

WPI; 2004-449666/42

Use of agent(s) that elevate intracellular cyclic adenosine monophosphate or intracellular calcium levels in neural tissue for modulating neurogenesis to treat central nervous system disorder.

Disclosure; SEQ ID NO 43; 77pp; English.

This invention relates to a novel method of modulating neurogenesis in the neural tissue of a patient exhibiting symptom(s) of a central nervous system disorder, such as neurodegenerative, ischaemic or learning and memory disorder or neurological trauma. The method involves at least one agent (A) that elevates intracellular cyclic adenosine monophosphate (CAMP) levels or at least one agent (B) that elevates intracellular Ca 2+ levels in the neural tissue, which is administered where (A) modulates and (B) induces neurogenesis. The invention may be useful for the production of compounds with a nootropic, neuroprotective, CNS-Gen, cerebroprotective, vasotropic, anticonvulsant, antiparkinsonian. haemostatic, hypertensive, muscular-Gen, ophthalmological, antiinflammatory, analgesic or antidiabetic activity. These compounds may act as neurogenesis modulators, neural stream or progenitor cell proliferation, differentiation and/or migration modulators, neural tissue G-protein coupled receptor activators, neurogenesis inducers, intracellular neural cAMP enhancers, intracellular neural cAMP stimulators or intracellular neural Ca 2+ enhancers. The inventi useful for modulating neurogenesis in neural tissue of a patient exhibiting at least one symptom of central nervous system disord The invention nervous such

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cc as Parkinson's disease and Parkinson's disorders, Huntington's disease, can Alzheimer's disease, multiple sclerosis, amyotrophic lateral sclerosis, companies of the professive supranuclear palsy, Lewy body disease, spinal ischaemia, ischaemic stroke, cerebral infarction, spinal cord conjury, cancer-related brain and spinal cord injury, multi-infarct comentia and geriatric dementia; for increasing the intracellular levels of or stimulating cAMP levels in a cell (preferably a cell from a neural tissue); and for in vitro modulation of neurogenesis. The agent modulates neurogenesis in neural tissue by modulating proliferation, cells in the tissue; by maintaining or increasing the amount or cells in the tissue; by maintaining or increasing the amount or cells in the tissue; by maintaining or increasing the amount or coupled receptor in the neural tissue cells in the neural tissue relative to a patient not dosed with the agent or by activation of a G-protein coupled receptor in the neural tissue. The method results in elevation of campled receptor in the neural tissue. The method results in elevation of disorders caused by cell loss, injury or disease by endogenous cell sorters caused by cell loss, injury or disease by endogenous or patient. Neurogenesis can also be induced by administration of the neurogenesis allows treatment of cell suncecessary systemic administration and possible side effects and further provides an alternative to the use of drugs and the controversial use of large quantities of embryonic tissue for treatment of Parkinson's cell shown to have the ability to increase intracellular cAMP levels and which the invention.
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Sequence 52 AA,

Matches Query Match Local 15; Similarity Conservative 28.5%; 38.5%; 9 Pred. Mismatches 72.5; No. 0. . 028 DB 8; 17; Length Indels 52 1; Gaps 1;

밁 δ 15 9 GCRFGTCTVQKLAHQIYQFT-DKDKDNVAPRSKISPQGY GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47 52

ADP18402 standard; protein; 52

26-AUG-2004 (first entry

Neurogenesis modulation-related protein SeqID16

RESULT 14
ADP18402
ID ADP18
XX ADP18
XX ADP18
XX ADP18
XX ADP18
XX ADP18
XX Neuro
AX RW neurodegenerative; ischaemic; learning and memory disorder;

KW neurological trauma; nootropic; neuroprotective; CNS-Gen;

KW derebroprotective; vasotropic; anticonvulsant; antiparkinsonian;

KW hemostatic; hypertensive; muscular-Gen; ophthalmological;

KW hemostatic; hypertensive; muscular-Gen; ophthalmological;

KW neural stem cell; progenitor cell;

KW neural tissue G-protein coupled receptor activator; neurogenesis inducer;

KW intracellular neural cAMP enhancer; intracellular neural cAMP stimulator;

KW intracellular neural Ca 2+ enhancer; parkinson's disease;

KW intracellular neural Ca 2+ enhancer; Parkinson's disease;

KW multiple sclerosis; amyotrophic lateral sclerosis; Shy-Drager syndrome;

KW progressive supranuclear palsy; Lewy body disease; spinal ischaemia;

KW schaemic stroke; cerebral infarction; spinal cord injury;

W cancer-related brain; spinal cord injury; multi-infarct dementia;

KW geriatric dementia; cAMP level; embryonic tissue; human. neurogenesis modulation; neural tissue; central nervous system disorder;

03-JUN-2004.

20-NOV-2003; 2003WO-IB005311

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CC cerebroprotective, vasotropic, anticonvulsant, antiparkinsonian.

CC haemostatic, hypertensive, muscular-Gen, ophthalmological,

antiinflammatory, analgesic or antidiabetic activity. These compounds may

CC antiinflammatory, analgesic or antidiabetic activity. These compounds may

CC antiinflammatory, analgesic or antidiabetic activity. These compounds may

CC act as neurogenesis modulators, neural stem or progenitor cell

CC proiferation, differentiation and/or migration modulators, neural tissue

CC groiferation, differentiation and/or migration modulators, neural tissue

CC intracellular neural cAMP enhancers, intracellular neural cAMP

CC stimulators or intracellular neural C2 + enhancers. The invention is

CC stimulators or intracellular neural case of a patient

CC exhibiting at least one symptom of central nervous system disorder, such

CC as Parkinson's disease and Parkinson's disorders, Huntington's disease,

CC shy-Drager syndrome, progressive supranuclear palsy, Lewy body disease,

CC shy-Drager syndrome, progressive supranuclear palsy, Lewy body disease,

CC shipury, cancer-related brain and spinal cord injury, multi-infarct

CC dementia and geriatric dementia; for increasing the intracellular levels

CC of or stimulating cAMP levels in a cell (preferably a cell from a neural

CC tissue); and for in vitro modulation of neurogenesis. The agent modulates

CC differentiation, migration or survival of neural stem cells or progenitor

CC cells in the tissue; by maintaining or increasing the amount or

CC percentage of doublecortin positive cells in the neural tissue relative

CC or patient mot dossed with the agent or by activation of a G-prortein
to a patient not dosed with the agent or by activation of a G-protein coupled receptor in the neural tissue. The method results in elevation CAMP levels of the neural cells by one method results to untreated tissue. The in vivo induction of neurogenesis allows treatment of disorders caused by cell loss, injury or disease by endogenous replacement and obviates the need for transplanting foreign cells into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the neural tissue of a patient exhibiting symptom(s) of a central nervous system disorder, such as neurodegenerative, ischaemic or learning and memory disorder or neurological trauma. The method involves at least one agent (A) that elevates intracellular cyclic adenosine monophosphate (cAMM) levels or at least one agent (B) the neural tissue, which is administered where (A) modulates and (B) induces neurogenesis. The invention may be useful for the production of compounds with a nootropic, neuroprotective, CNS-Gen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Use of agent(s) that elevate intracellular cyclic adenosine monophosphate or intracellular calcium levels in neural tissue for modulating neurogenesis to treat central nervous system disorder.
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Roennholm H, Zach
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MERCER A.
NJBERG K.
OSSOINAK A.
PATRONE C.
ROENNHOLM H.
ZACHRISSON O.
WIKSTROEM L.
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Haeggblad J, Jansson A,
Mercer A, Njberg K,
O, Wikstroem L;
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Best Local
                                                                           The present invention relates to a method for identifying drug non-target biomolecules in a mixture of biomolecules. The method comprises interacting mixture with capture compounds having moiety X which covalently binds to biomolecules with high affinity, moiety Y that increases selectivity of binding so that the capture compound binds to fewer biomolecules, and moiety Z for presenting X and Y, and analysing captured biomolecules to identify drug non-targets. The capture compound also optionally comprises a sorting function moiety Q and or a solubility function moiety W. The selectivity function moiety Y serves to modulate
                            the reactivity function by reducing the number of groups to which the reactivity function moiety X bind, such as by steric hindrance and ot interactions. Y is optionally a peptide ligand (ADR42112-ADR42256).
                                                                                                                                                                                                                                                                       Identifying drug non-target biomolecules in mixture of biomolecules involves interacting mixture of biomolecules with capture compounds having high binding affinity and analyzing captured biomolecules to identify drug non-targets.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; ligand; Adrenomedullin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neurogenesis-modulating agent directly into a desired site, which avoids unnecessary systemic administration and possible side effects and further provides an alternative to the use of drugs and the controversial use of large quantities of embryonic tissue for treatment of Parkinson's disease. The present sequence is that of a polypeptide which has been shown to have the ability to increase intracellular cAMP levels and which is related to the method of the invention.
                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 2; 368pp; English
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Hassman CF,
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Yip P;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAMP-dependent pathway.
SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Expressed in the esophagus,
ileum, ileocecum, ascending colon, transverse co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FUNCTION: IMDL and IMDS may play a role as physiological regulators of gastrointestinal, cardiovascular bioactivities
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SIMILARITY: Belongs to the adrenomedullin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mediated by the CALCRL/RAMPs receptor complexes. Activates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           608682;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SWISS-PROT entry is copyright. It is produced through sen the Swiss Institute of Bioinformatics and the EN
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                                                                                                                                                                                                                                                                                                                           47;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HGNC:28898; ADM2.
                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                      TQAQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                                                                                            TQAQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 147
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44, Last annotation update)
precursor (Intermedin) (Cont.
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                                   Intermedin-short
                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of basic residues;
                                                                                                                                                                                                                                                                                                                                             Score
Pred. 1
                                                                                                                                                                                                                                                                                                                                                                                                                              By similarity.
Adrenomedullin 2 (By similarity Intermedin-short (Potential).
By similarity.
By similarity.
Tyrosine amide (G-148 provides
                                                                                                                                                                                                                                                                                                                                                                                                                     group)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Potential.
                                                                                                                                                    PRT;
                                                                                                                                                                                                                                                                                                                                                                                                     6E0E3098CFCE5BF2 CRC64;
                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                           254; L-
No. 1.2e-24;
                                                                                                                                                                                                                                                                                                                                                                                                                        (Probable)
                                                                                                                                                    150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      moved. Usage by and for commercial (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          There are no restrictions ong as its content is in
                                                     (Contains:
                                                                                                                                                  A
                                   (IMDS)].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hormone; Signal.
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                                                                                                                                                                                                                                                                                                                                                           Length 148;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            similarity).
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                                                       Adrenomedullin
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Best Local
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P61312;
05-JUL-2004
05-JUL-2004
05-JUL-2004
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the Euro
                   05-JUL-2004 (Rel. 44, Created)
05-JUL-2004 (Rel. 44, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Adrenomedullin 2 precursor (Intermedin) (Cont.
                                                                                                                                                                                                                                                                                                 BEPTIDE
                                                                                                                                                                                                                                                                                                                                                                         EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Roh J., Chang C.L., Bhalla A., Klein C., Hsu S.Y.T., "Intermedin is a calcitonin/calcitonin gene-related peptide acting through the calcitonin receptor-like activity-modifying protein receptor complexes.", J. Biol. Chem. 279:7264-7274 (2004).
                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                              entities requires a license agreement (S or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                      modified
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                      SIGNAL
                                                                                                                                                                                                                                                                                                                                                               MGD; MGI:2675256;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PubMed=14706825;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=Adm2; Synonyms=Am2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mediated by concentrations and mediated by the submaxillary SUBCELLULAR LOCATION: Secreted.

SUBCELLULAR LOCATION: Secreted.

TISSUE SPECIFICITY: High expression detected in the submaxillary rissue specification, and mesentery, followed by the pituitary, reland, kidney, stomach, and mesentery, followed by the pituitary, reland, kidney, stomach, and mesentery, followed by the pituitary, with sporadic i
                                                                                                                                                                                                                                                                                                                                                                                                                                                             s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation -
European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SIMILARITY: Belongs to the adrenomedullin family.
                                                                                                                                                  105
                                                                                                                                                                                                                                                                                                                                                                            AB121035;
                                                                                                                                                                           ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             anterior lobe.
                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                           non-profit institutions as long as its content and this statement is not removed. Usage by an requires a license arenous temoved.
                                                                                                                                                AQLLRVGCVLGTCQVQNLSHRLWQLVRPAGRRDSAPVDPSSPHSY 149
                                                                                                                                                                    AQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                                                                                                  150
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103
110
112
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BAD07412.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DOI=10.1074/jbc.M305332200;
L., Bhalla A., Klein C., Hsu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DOI=10.1016/S0014-5793(03)01368-1;
                                                                                                                                                                                                                                                                                                                                                                Adm2.
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117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chordata;
                                                                                                                                                                                                            89.8%;
                                                                                                                                                                                                                                                                                                                                                  pair
                                                                                                                                                                                                                                                  M.
                                                                                                                                                                                                 Score 228; DB Pred. No. 2.6e 2; Mismatches
                                                                                                                                                                                                                                                                         By similarity.
Adrenomedullin 2
Intermedin-short
By similarity.
Tyrosine amide (G-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AND TISSUE SPECIFICITY
                                                                                                                                                                                                                                                                                                                                                   basic
                                                                                                                                                                                                                                                  oup) (Probable).
2BFF392BFBFDEBBA
                                                                                    146
                                                                                                                                                                                                                                                                                                                                                  residues; Hormone; Signal
                                                                                                                                                                                                                                                                         amide (G-150 provides amide
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(Potential).
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receptor/receptor
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(Intermedin-long)

Intermedin-short (IMDS)].

(Contains:

Adrenomedullin

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Best Local S
Matches 41
                                                                                                             Q75XW6;
Q75XW6;
05-JUL-2004
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DISULFID
MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                          the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fugu rubripes (Japanese pufferfish) (Takifugu rubripes).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterostei; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Takifugu.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PubMed=14706825; DOI=10.1016/S0014-5793(03)01368-1; Takei Y., Inoue K., Ogoshi M., Kawahara T., Bannai H., "Identification of novel addrenomedullin in mammals: a cardiovascular and renal regulator."; PEBS Lett. 556:53-58(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                        EMBL;
                                                                   Name=ADM3;
                                                                                        05-JUL-2004
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                                                                              Adrenomedullin-3.
                                                                                                                                                         Δ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SIMILARITY: Belongs to the adrenomedullin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Expression was restricted to and anterior lobes of the pituitary.
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                                                                                                                                                                                                                                                                                                                                                                                        AB121036; BAD07413.1;
                                                                                                                                                                                                                 w
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                                                                                                                                                                                          AQLLRVGCVLGTCQVQNLSHRLWQLVRPSGRRDSAPVDPSSPHSY
                                                                                                                                                                                                          AQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
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91.1%;
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Pred. No. 6.2e-21;
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3; Mismatches
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                                                                                                                                                                                                                                                                                   group) (Probable).
C87043237AD29DDC
                                                                                                                                                                                                                                                                                                                                          By similarity.
Adrenomedullin 2
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                                                                                                                                    PRT;
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                                                                                                                                      159
                                                                                                                                                                                                                                                                                                                                                                           residues; Hormone; Signal.
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                                                                                                                                                                                                                                                            Length 146
                                                                                                                                                                                                                                                                                    CRC64;
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Best Local
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Eukaryota; Metazoa; Choruaca,
Actinopterygii; Neopterygii;
Currinidae; Danio
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Q75XW7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fugu rubripes (Japanese pufferfish) (Takifugu rubripes).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Actinopterygii; Neoptergogii; Teleostei; Buteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;

Tetradontoidea; Tetraodontidae; Takifugu.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.

MEDLINE=22984567; PubMed=14623291; DOI=10.1016/j.bbrc.2003.10.111;

Ogoshi M., Inoue K., Takei Y.;

"Identification of a novel adrenomedullin gene family in teleost
PubMed=14615490; DO Roh J., Chang C.L., "Intermedin is a ca
                                                                                                                                                                 Brachydanio rerio (Zebrafish)
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                   05-JUL-2004
05-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AB120296; BAD02342.1; -.
InterPro; IPR001938; Thaumatin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. MEDLINE=22984567;
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05-JUL-2004
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                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                              Name=IMDN;
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                                                                                                                                                                                                                                                                                                                 Q6IFS9;
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Identification of a novel adrenomedullin gene family
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  DOI=10.1074/jbc.M305332200;
L., Bhalla A., Klein C., Hsu calcitonin/CGRP family pepti
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Pred. No. 4e-16;
7; Mismatches
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Pred. No. 5.6e-16;
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RESULT
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                                     Fugu rubripes (Japanese pufferfish) (Takifugu rubripes).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoldea; Tetraodontidae; Takifugu.
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Q75XW8;
05-JUL-2004
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NON TER
SEQUENCE
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05-JUL-2004 (TrEMBLrel.
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Submitted
                      NCBI_TaxID=31033;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Takifugu.
                                                                                                                                                     Name=ADM5;
                                                                                                                                                                         Adrenomedullin-5.
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-!- MISCELLANEOUS: The sequence shown here is derived from EMBL/GenBank/DDBJ third party annotation (TPA) entry.
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Identification of a novel adrenomedullin
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GO:0005576; C:extracellular; IEA.
GO:0005179; F:hormone activity; IEA.
GO:0005179; F:hormone activity; IEA.
UENCE 174 AA; 20222 MW; 61535E41FCF88D4D CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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Pred. No. 0
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Pred. No. 7.
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01-FEB-1994 (Rel. 2
01-FEB-1994 (Rel. 2
05-JUL-2004 (Rel. 4
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SIGNAL
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                                                                                                   01-FEB-1994 (Rel. 28, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
ADM precursor (Contains: Adrenomedullin (AM);
terminal peptide (ProAM-N20) (ProAM N-terminal
                      Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                 Homo sapiens (Human)
                                                                                          Name=ADM;
                                                                                                                                                                                                                                            ADML
                                                                                                                                                                                                                                                                   HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q6L8K5;
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MEDLINE=22984567; PubMed=14623291; DOI=10.1016/j.bbrc.2003.10.111;
Ogoshi M., Inoue K., Takei Y.;
"Identification of a novel adrenomedullin gene family in teleost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CHAIN
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Gen. Comp. Endocrinol. 138:78-88(2004)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Molecular cloning and expression of preproadrenomedullin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PubMed=15242754; DOI=10.1016/j.ygcen.2004.05.002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Name=preproAM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CISSUE=Liver;
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L; AB120940; EAD19046.1; -.
G0:0005576; C:extracellular; IEA.
G0:0005179; F:hormone activity; IEA.
                                                                                                                                                                                                                                            HUMAN
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                                                                                                                                                                                                                                                                                                                                                      SQSRRSGCSLGTCTVHVLAHRLHDLNNKL-KIGNAPADKINPFGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GCQVGTCQVHNLANKLYQIGQRQGKDESTKV--NDPQGY
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                                                                                       Synonyms=AM;
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79 127 a
171 AA; 19412 MW;
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                                                                                                                                                                                                                                            STANDARD;
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                      Chordata;
Primates;
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Pred. No.
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Pred. No. 0.053;
5; Mismatches
                      Craniata; V
Catarrhini;
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i; Hominidae;
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l 20 peptide) (PAMP)].
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RA Strausberg R.L., Peingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altechul S.F., Zeeberg B., Buctow K.H., Schaefer C.F., Bhat N.K.,
RA Altechul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Moßwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Raha S.S., Moßwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Roberds S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Mozny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Hakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
ROdriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J.M., Marra M.A.;
Tonco Matl A. Rodrigues S., Sanchez B.,
The C. Matl A. Rodrigues S.,
RA Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.",
                                                                                                                                                                                                                 Regul.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                +
                                                                                                                                                                                                                                                                      MEDLINE=20053666; PubMed=10588445; DOI=10.1016/S0167-0115(99)00025-7; Champion H.C., Nussdorfer G.G., Kadowitz P.J.; "Structure-activity relationships of adrenomedullin in the circulation
                                                                                                                                                                                                                                                                                                                                                                                "Proadrenomedullin-derived peptides."; Front. Neuroendocrinol. 19:100-127(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                            Samson W.K.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=98240137;    PubMed=9578982;    DOI=10.1006/frne.1998.0164;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Adrenomedullin: a novel hypotensive peptide isolated from human pheochromocytoma.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matsuo H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Pheochromocytoma;
MEDLINE=93343928; PubMed=7688224;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                iochem.
                                 FUNCTION: AM and PAMP are potent hypotensive and vasodilatator agents. Numerous actions have been reported most related to the physiologic control of fluid and electrolyte homeostasis. In the kidney, am is diurettc and natriurettc, and both am and pamp inhibit aldosterone secretion by direct adrenal actions. In pituitary gland, both peptides at physiologically relevant doses inhibit basal ACTH secretion. Both peptides appear to act in brai and pituitary gland to facilitate the loss of plasma volume, actions which complement their hypotensive effects in blood
                                                                                                                                                                                                                              adrenal gland.";
ul. Pept. 85:1-8(1999).
SUBCELLULAR LOCATION: Secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Biophys. Res. Commun.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eto T.;
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Best Local S
Matches 15
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GO; GO:0005615; C:extracellular space; TAS.
GO; GO:0005625; C:soluble fraction; TAS.
GO; GO:0005102; F:receptor binding; TAS.
GO; GO:0005171; P:cAMP biosynthesis; TAS.
GO; GO:0006171; P:cAMP biosynthesis; TAS.
GO; GO:0007267; P:cell-cell signaling; TAS.
GO; GO:0008015; P:circulation; TAS.
GO; GO:0007565; P:pregnancy; TAS.
GO; GO:0007565; P:pregnancy; TAS.
GO; GO:0005701; P:progesterone biosynthesis;
GO; GO:0009611; P:response to wounding; TAS.
GO; GO:0007165; P:signal transduction; TAS.
InterPro; IPR001710; Adrenomedullin.
                                                       ADML POUR POTAS; P97297; P97453; P97297; P97453; P67297; P97453; P67297; P97453; P67297; P672901 (Rel. 40, Last sequence update) 16-OCT-2001 (Rel. 44, Last annotation update) 05-JUL-2004 (Rel. 44, Last annotation update) ADM precursor [Contains: Adrenomedullin (AM); P67441 Deptide (ProAM-N20) (ProAM N-terminal)
                                                                                                                                                                                        MOUSE
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DISULFID
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NCBI_TaxID=10090;
                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                         VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                    MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                             MOD_RES
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Genew; HGNC:259; ADM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; BC015961; AAH15961.1;
EMBL; D43639; BAA07756.1; A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; D14874; BAA03589.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -i- TISSUE SPECIFICITY: Highest levels found in pheochromocytoma and
adrenal medulla. Also found in lung, ventricle and kidney tissues
-i- SIMILARITY: Belongs to the adrenomedullin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amidation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRINTS; PR00801; ADRENOMEDULN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF02039; Adrenomedullin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     H-InvDB; HIX0009441; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein
                                                                                                                                                                                                                                                                        9 GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                            AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequencing; Hormone; Polymorphism; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P:progesterone biosynthesis;
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185
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38.5%;
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ALT_SEQ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of basic residues;
                                                                                                                                                                                                                                                                                                                 Score 72.5; DB Pred. No. 0.34;
                                                                                                                                                                                                                                                                                                                                                                          /FTId=VAR_014861
                                                                                                                                                                                                                                                                                                                                                                                                                  Tyrosine amide (G-147 provides amide
                                                                                                                                                                                                                                                                                                                                                                                                                                               Arginine amide (G-42 provides amide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PreproAM C-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Adrenomedullin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Proadrenomedullin N-20 terminal peptide
                                                                                                                                                                                                                                                                                                                                                              64C7D2A0B4654DFE CRC64;
                                                                                                                                                                                                                                                                                                                                                                                         -> R (in dbSNP:5005).
                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ventricle and kidney tissues.
                                                                                                                                                                                                                                                                                                                              DB 1; Length 185;
                                                                                                                                                                                                                                                                                                     17;
                                                                                                  Proadrenomedullin N-20
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Best Local
                                                                                            ADML_BOVIN
062827;
16-0CT-2001
                                                                                                                                                                                                                                                                        CONFLICT
                                               062827;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
05-JUL-2004 (Rel. 44, Last annotation update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                               BOVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        trophoblast giant cells at the embryo implantation sidev. Biol. 203:264-275(1998).
 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                              terminal peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through a collaboratic between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mouse adrenomedullin gene.";
Genomics 37:395-399(1996).
                        Bos taurus (Bovine)
                                    Name=ADM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                    SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                               Amidation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR001710; Adrenomedullin.
Pfam; PF02039; Adrenomedullin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MGD; MGI:108058; Adm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; U77630; AAB36535.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; D78349; BAA11367.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KO M.S.H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yotsumoto S., Shimada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE-99046755; PubMed-9808778; DOI=10.1006/dbio.1998.9073;
Yotsumoto S., Shimada T., Cui C.Y., Nakashima H., Fujiwara H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                       Local
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SIMILARITY: Belongs to the adrenomedullin family.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               non-profit institutions as long and this statement is not removed.
                                                                                                                                                                              GCRFGTCTFQKLAHQIYQLT-DKDKDGMAPRNKISPQGY
                                                                                                                                                                                                    GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                                                                                                                                                                                                                                                                                                                                                                              Cleavage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         equires a license agreement (S email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                         184 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92892; PubMed=8938454; DOI=10.1006/geno.1996.0576; Ogawa Y., Tamura N., Mori Y., Isse N., Aoki T., M., Taketo M.M., Seldin M.F., Nakao K.;
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 Cetartiodactyla;
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             Chordata;
                                                                                                                                                                                                                                    28.1%;
38.5%;
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                                                                                                                                                                                                                                                                         MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                           of basic residues; Hormone; Signal.
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                                                                                                                                                                                                                          Score 71.5; D
Pred. No. 0.46
6; Mismatches
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A -> G
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PreproAM C-terminal fragment similarity).
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                                                                                                                                                                                                                                                                                                                                                                                                     By similarity. Proadrenomedullin N-20 terminal peptide.
                                                                                                                                                                                                                                                                                                                                              By similarity.
                                                                                                                                                                                                                                                                                                                                                                                            By similarity.
            Craniata; Vertebrata; Euteleostomi;
                                                                                                                    PRT;
                                                                                                                                                                                                                                                                         C88C99045A79C898
                                               N-terminal
                                                                                                                                                                                                                                                                                    (By similarity). (in Ref. 2).
 Ruminantia; Pecora;
                                                                                                                    188
                                                                                                                                                                                                                                                                                                         amide (G-145 provides amide
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                                             Proadrenomedullin N-20 l 20 peptide) (PAMP)].
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 Bovidae;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                          Q95KP0;
Q95KP0;
01-DEC-2001
                                                                                      SEQUENCE FROM N.A.
MEDLINE=21630318; PubMed=11754956; DOI=10.1016/S0196-9781(01)00529-0;
Viramura K.. Matsui E., Kato J., Katoh F., Kita T., Tsuji T.,
                                                                                                                                                                                NCBI_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DISULFID MOD_RES
               Kangawa K., Eto T.;
"Adrenomedullin (11-26): a novel endoge isolated from bovine adrenal medulla.";
Peptides 22:1713-1718 (2001).
                                                                                                                                                                                                               Mammalia; Butheria;
Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                   01-DEC-2001 (TrEMBLrel. 19,
01-DEC-2001 (TrEMBLrel. 19,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce) or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Cloning of bovine preproadrenomedullin and inhibition of expression in vascular endothelial cells by staurosporine. Life Sci. 62:1407-1415(1998)
-I- FUNCTION: Hypotensive peptide. May function as a hormoned the sci. 62:1407-1415(1998)
                                                                                                                                                                                                                                                   Bos taurus (Bovine).
Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                               Name=pBAM-2;
                                                                                                                                                                                                                                                                                                                  Adrenomedullin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9913;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EPTIDE
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Buropean Bioinformatics Institute. There are no rest
by non-profit institutions as long as its content
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  circulation
                                                                                                                                                                                         TaxID=9913;
    AB055107;
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                                                                                                                                                                                                                              Chordata; Craniat
Cetartiodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clark A.J.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20981 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28.1%;
38.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 71.5; DB 1;
Pred. No. 0.47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   group) (B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            group)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            By similarity. Arginine amide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PreproAM C-terminal similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                By similarity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Proadrenomedullin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         By similarity
                                                                                                                                                                                                                                                   Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3002E79AB3B6612C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                            endogenous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Corder R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (By similarity).
ne amide (G-147 provides
(By similarity).
                                                                                                                                                                                                                                  Ruminantia; Pecora;
                                                                                                                                                                                                                                                                                                                                                                                                                       188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            residues;
                                                                                                                                                                                                                                                                                                                                                                                                                       B
                                                                                                                                                                                                                                                                                                                                                           update)
                                                              hypertensive peptide
                                                                                                                                                                                                                                                                                                                                   update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-20 terminal peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hormone; Signal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      188;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL
                                                                                                                                                                                                                                  Bovidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              its.
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P53366;
01-OCT-1996
01-OCT-1996
                                                                                           SIGNAL
PEPTIDE
                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sbb.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=94357274; PubMed=8076689; DOI=10.1016/0014-5793(94)00810-8; Kitamura K., Kangawa K., Ishiyama Y., Washimine H., Ichiki Y., Kawamoto M., Minamino N., Matsuo H., Eto T.; Rawamoto M., Minamino N. Matsuo H., Eto T.; "Identification and hypotensive activity of proadrenomedullin N-terminal 20 peptide (PAMP)."; FEBS Lett. 351:35-37(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GO; GO:0005576; C:extracellular; IEA.
GO; GO:0005179; F:hormone activity; I
InterPro; IPR001710; Adrenomedullin.
Pfam; PF02039; Adrenomedullin; 1.
PRINTS; PR00801; ADRENOMEDULN.
                                                                                                                                                                                                                                                                                                                             <del>;</del>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FAVE. ......
TISSUE=Addrenal medulla;
MEDLINE=94139945; PubMed=8043068; DOI=10.1016/0014-5793(94)80289-0;
Kitamura K., Kangawa K., Kojima M., Ichiki Y., Matsuo H., Eto T.;
"Commolete amino acid sequence of porcine adrenomedullin and cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
                                        Direct protein
                                                                                                                                                                                                                                                                                              -!- SIMILARITY: Belongs to the adrenomedullin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sus scrofa (Pig)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     terminal peptide (ProAM-N20) (ProAM N-terminal 20 peptide) (PAMP)].
Name=ADM; Synonyms=AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ol-OCT-1996 (Rel. 34, Created)
Ol-OCT-1996 (Rel. 34, Last sequence update)
O5-UL-2004 (Rel. 44, Last annotation update)
ADM precursor [Contains: Adrenomedullin (AM);
                                                                       PRINTS; PR00801; ADRENOMEDULN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CISSUE=Adrenal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE OF 22-41
                                                                                                                                                                                                                                                                                                                          SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Highly expi
                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: AM and PAMP are potent hypotensive and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lett.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        109
                                                                                     PF02039; Adrenomedullin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9 GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCRFGTCTVQKLAHQIYHFT-DKDKDGSAPRSKISPQGY 146
                                                       Cleavage on pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          188 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
22
45
                                         sequencing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               medulla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Suina; Suidae; Sus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20963 MW;
                                         Hormone;
                                                         of basic residues;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5
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; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 71.5;
            Proadrenomedullin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6102E69A756DCA86 CRC64;
                                                                                                                                                                                                                                                                                                                            expressed in adrenal glands,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No. 0.47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IEA.
                                         Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             188
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Proadrenomedullin N-20
             N-20
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            terminal
                                                                                                                                                                                                                                                                                                                                                                         vasodilatator
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                                                                                                                                                                                                                                                                                                                          lung
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          peptide.
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                                                        Query Match
Best Local S
Matches 13
                                                                                                                                                                                                                                                                                      Galagan J.E., Calvo S.E., Borkovich K.A., Selker B.U., Read N.D.,
Jaffe D., FitzHugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B.,
Ra Elkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M.,
Qui D., Ianakiev P., Pedersen D., Nelson M., Washburne M.,
Qui D., Ianakiev P., Pedersen D., Nelson M., Washburne M.,
Qui D., Ianakiev P., Pedersen D., Nelson M., Washburne M.,
Roy L., Foldd G., Mewes W., Staben C., Marcotte E., Greenberg D.,
Roy A., Foley K., Naylor J., Thomann N., Barrett R., Gnerre S.,
Roy A., Foley K., Naylor J., Thomann N., Barrett R., Gnerre S.,
Roy A., Foley K., Naylor J., Mauceli E., Bielke C., Rudd S., Frishman D.,
R. Kamal M., Kamvysselis M., Mauceli E., Bielke C., Rudd S., Frishman D.,
R. Kamal M., Kamvysselis M., Mauceli E., Bielke C., Rudd S., Frishman D.,
R. Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S.,
A. Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S.,
A. Cogoni C., Macino G., Catcheside D., Li W., Pratt R.J., Osmani S.A.,
A. DeSouza C.C., Glass L., Orbach M.J., Berglund J., Voelker R.,
A. Natvig D.O., Alex L.A., Mannhaupt G., Ebbole D.J., Freitag M.,
Rabert R., Sache M.S., Lander E.S., Nusbaum C., Birren B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
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Q7S575;
Q1-MAR-2004
Q1-MAR-2004
Q1-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neurospora crassa.
Eukaryota; Fungi; Ascomycota; Pe
Sordariomycetidae; Sordariales;
                                                                                                                 Hypothetical SEQUENCE 97
                                                                                                                                        InterPro; IPR002498; Fir: Pfam; PF01504; PIP5K; 1.
                                                                                                                                                                                                                              -!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry
                                                                                                                                                                                                                                                              Nature 0:0-0(2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=OR74A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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                                                                                                                                                                                          HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=5141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=NCU02295.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hypothetical protein.
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                                                                                                                                                                                                                                                                           "The Genome Sequence of the Filamentous Fungus Neurospora crassa.";
                                                                                                                                                          iP; P78356; 1BO1.
GO:0016308; F:1-phosphatidylinositol-4-phosphate 5-kinase.
serPro; IPR002498; PIP5K.
                                                                                                                                                                                                    preliminary data.; AABX01000355; E
 123
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                            30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9 ĠĊVLĠŢĊQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY
                                                                       Similarity
PAQRRDSAPSSPSSPHRY
                            PAGRODSAPVDPSSPHSY 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCRFGTCTVQKLAHQIYQFT-DKDKDGVAPRSKISPQGY 146
                                                                                                                   927 AA; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 188 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TrEMBLrel. (TrEMBLrel.
                                                         Conservative
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153
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41
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188
                                                                                                                102558 MW;
                                                                                                                                                                                                      EAA30703.1;
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38.5%;
                                                                       27.6%;
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Pred. No. 0.63
5; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Created)
                                                                       Score 70;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               group).
71749460F5660A61 CRC64;
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Tyrosine amide (G-147 provides amide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         By similarity. Arginine amide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Adrenomedullin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pezizomycotina; Sordariomycetes; s; Sordariaceae; Neurospora.
                                                           Mismatches
                                                                                                                 21FDC49FA419E932 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           927
                                                                                    DB 2; Length 927;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Selker E.U., Re
S., Purcell S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (G-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1;
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                                                         4.
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                                                         Indels
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Search completed: May Job time : 178 secs

18:14:12

152657 A051657 A0516555 G711409 E8513409 E851357 T10235 T102392 T102392 T102392 T102392 T12022 T12022 T12022 T12022 T15022 T15022 T15022 T15022 T15022

xanthine dehydroge genome polyprotein alpha-2-macroglobu hypothetical prote peroxidase (EC 1.1 hypothetical prote hypothetical prote

407 552 771 813 1359 3898 4544 296 296 317 317 3460 488 503

50.5 50.5 50.5 50.5

hypothetical reduced 12K membrane glyco NAD synthase (EC 6 probable replication hypothetical protestion)

seizure-related pr hypothetical reduc

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Result
No.
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                Post-processing: Minimum Match 0%
                                                                                                                                                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                        Searched:
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                                                                       d. No. is the number of results predicted by chance to have a
re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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Match
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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Listing first 45 summaries
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115.954 Million cell updates/sec
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ALIGNMENTS

ative (	obable (EC 1.1.1
Best Local Similarity 38.5%; Pred. No. 0.037;	lyprotein - hog
	nome polyprotein
F;146/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following gl)	clic beta 1-2 gl
F. 110-115 Dismifide bonds: #Fratus experimental	potmeticai proce obable integral
F:147-185/Domain: Carboxyl-terminal propeptide #status predicted cCTPD-	notbetical prote
F;95-146/Product: adrenomedullin #status experimental <mat></mat>	pothetical prote
F;22-41/Domain: proadrenomedullin amino-terminal 20 peptide #status predicted <pap></pap>	bB kinase activa
F;22-185/Product: proadrenomedullin #status predicted <peu></peu>	bB kinase activa
F;1-21/Domain: signal sequence #status predicted <sig></sig>	pothetical prote
C:Keyords: amidated carboxyl end: blood pressure control: hormone	steine proteinas
A:Introne: 3J/2: 8J/C	steine proteinas
A.Man Docition   11ster   1ster   Crimitals	TD-D-Glucose 4 6
A;Gene: GUB:AUM	pocnecicai proce
C;Genetics:	pothetical prote
A;Experimental source: pheochromocytoma	rboxylesterase (
A;Residues: 95-146 <ki3></ki3>	mma-interferon-i
A; Molecule type: protein	pothetical prote
A;Accession: JN0476	C transporter AT
A; Reference number: JN0476; MUID: 93249425; PMID: 8387282	c16-related prot
A; Title: Adrenomedullin: A novel hypotensive peptide isolated from human pheochromocytoma	pothetical prote
Biochem. Biophys. Res. Commun. 192, 553-560, 1993	pha-1A adrenergi
R; Kitamura, K.; Kangawa, K.; Kawamoto, M.; Ichiki, Y.; Nakamura, S.; Matsuo, H.; Eto, T.	lactate permease
A; Residues: 22-41 < KI2>	renomedullin pre
A; Molecule type: protein	renomedullin - p
A:Accession: PN0548	renomedullin pre
A:Cross references GB:D14874: NID:G455470: DIDN:BAA03589 1: DID:G500612	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
A.Radidinas 1-185 KTTV	scription
A;ACCESSIDI: UNUBUS A:MOIDO: DENBA	
A; Reference number: JN0684; MUID:93343928; PMID:7688224	
A; Title: Cloning and characterization of cDNA encoding a precursor for human adrenomedul:	
Biochem. Biophys. Res. Commun. 194, 720-725, 1993	ion.
R; Kitamura, K.; Sakata, J.; Kangawa, K.; Kojima, M.; Matsuo, H.; Eto, T.	being printed,
A: Experimental source: pheorical conservations	to have a
OFFRENCHIO : CASUSOTTATION OFFRENCHION OFFRENCHION OF TRANSPORTED THE SECURITY SECUR	
A; MOLECULE TYPE: UNA	
A; Accession: JC2351	
A;Reference number: JC2351; MUID:94354869; PMID:8074714	
A; Title: Genomic structure of human adrenomedullin gene.	
Biochem. Biophys. Res. Commun. 203, 631-639, 1994	
R; Ishimitsu, T.; Kojima, M.; Kangawa, K.; Hino, J.; Matsuoka, H.; Kitamura, K.; Eto, T.;	
C:Accession: JC2351; JN0684; JN0476	
C.Date: Now begress was:	
Grenomeduliin precursor - numan	

222 222

\$12099 \$74598 JC5702 JC5701

T26050 A43708 A37845 T35335

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l-lactate permease [:
C;Species: Brucella r
C;Date: 01-Feb-2002 ;
C;Accession: AC3406
                                                                                                                                                                                                                                                                                                                                    C;Keywords: amidated carboxyl end
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-185/Product: proadrenomedullin #status predicted <PEU>
F;22-11/Product: proadrenomedullin amino-terminal 20 peptide #status predicted
F;24-143/Product: adrenomedullin #status predicted <AMI>
F;141/Modified site: amidated carboxyl end (Arg) (amide in mature form from foll
F;143/Modified site: amidated carboxyl end (Tyr) (amide in mature form from foll
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Blochem. Blophys. Res. Commun. 195, 921-927, 1993
4;Title: Molecular cloning and biological activities of
A;Reference number: JN0766; MUID:93384621; PMID:7690563
A;Accession: JN0766
                                                                                        AC3406
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FEBS Lett. 338, 306-310, 1994
A;Title: Complete amino acid sequence of porcine adrenomedullin and cloning
A;Reference number: S41600; MUID:9413945; PMID:8043068
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;Residues: 22-41 <SA2>;Comment: This precursor contains a unique 20-amino acid
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Species: Rattus norvegicus (Norway rat)
Date: 30-Sep-1993 #sequence_revision 20-Aug-1994 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Residues: 1-185 <SAK>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule type: mRNA;Residues: 1-188 <KIT>
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                                             [imported] - Brucella melitensis (strain
a melitensis
           #sequence_revision 01-Feb-2002 #text_change
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38.5%;
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Pred. No. 0.069;
5; Mismatches
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end (Tyr) (amide in mature form from following gly
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                                                        C;ACCEBBLOON, S.
R;Stellyes, L.; Gattung, S.
submitted to the EMBL Data Library, March 1996
submitted to the sequence of C. elegans cosm
                                                    A; Description: The sequence A; Reference number: Z20625
                                                                                                                                        C; Accession:
                                                                                                                                                   C:Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change
      A;Status: preliminary; translated
                                  A;Accession:
                                                                                                                                                                                                     hypothetical protein F08B4.2 - Caenorhabditis elegans
                                                                                                                                                                                                                                T29482
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Species: Caenorhabditis elegans

T29482

T29482

from

GB/EMBL/DDBJ

cosmid

F08B4

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A;Experimental source: hippocampus
A;Note: sequence extracted from NCBI backbone (NCBIP:73541)
C;Superfamily: vertebrate rhodopsin
C;Keywords: G protein-coupled receptor; glycoprotein; transmembrane
                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:P23944; GB:M60654; NID:g202761; PIDN: R;Perez, D.M.; Piascik, M.T.; Graham, R.M.
Mol. Pharmacol. 40, 876-883, 1991
A;Title: Solution-phase library screening for the identification A;Reference number: A53280; MUID:92100054; PMID:1661838
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                                              S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    alpha-1A adrenergic receptor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 22-Jan-1993 #text_change 09-Jul-2004
C;Accession: A38731; A53280
C;Accession: A38731; A53280
C;Accession: A38731; A53280
                                                                                                                                                                                                                                                                                            A;Accession: A53280
A;Status: preliminary; not compared with conceptual translation
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A; Residues: 1-560 < LOM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Lomasney, J.W.; Cotecchia, S.; Lorenz, W.; Leung, W.Y.; Schwinn, D.A.; Yang-Feng, T.L.; J. Biol. Chem. 266, 6365-6369, 1991
A;Title: Molecular cloning and expression of the cDNA for the alpha-lA-adrenergic receptor.
A;Reference number: A38731; MUID:91177889; PMID:1706716
                                                                                                                                                                                                                                                                        A;Residues: 1-36,'P',38-58,'I',60-202,'I',204-305,'R',307-366,'I',368-370,'I',372-559
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A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitens: A;Reference number: AD3252; PMID:11756688
A;Accession: AC3406
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT:Q8YGC6; GB:AB008917; PIDN:AAL52414.1; PID:g17983216; GSPDB:GN
A;Experimental source: strain 16M
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                                                                                              l Similarity
17; Conserv
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RLLR-----COCRRRRRLWSLRPPLASLDRRRAFRLRPQPSHRSPRGPSSPH
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                                                                                            Conservative
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                                                                                                          22.8%;
30.9%;
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27.7%;
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                                                                                                               Score 58; DB 2; Length 560 Pred. No. 10;
                                                                                       Pred. No. 10;
3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 58; DB :
Pred. No. 5.9;
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O'Callaghan, 1
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                                                                                     Gaps
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A; Molecule type: DNA
A; Residues: 1-283 <SIM>
A; Cooss-references: UNIPROT:Q9PD02; GB:AE003987; GB:AE003849; NID:g9106634; PIDN:AAF8441
A; Experimental source: strain 9a5c
A; Simpson, A.J.G.; Reinach, F.C.; Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R.; A
Briones, M.R. S.; Bueno, M.R. P.; Camargo, A.A.; Camargo, L.E.A.; Carraro, D.M.; Carrer, H
Briones, M.R. S.; Bueno, M.R. P.; Facincani, A.P.; Ferreira, A.J.S.
as-Neto, E.; Docena, C.; El-Dorry, H.; Facincani, A.P.; Ferreira, A.J.S.
submitted to GenBank, June 2000
A; Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco, M.C.; Frohn
A; Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.P.; Krieger, J.E.; Kuramae, E.E.; Laigy
Chado, M.A.; Madeira, A.M.B.N.; Madeira, H.M.F.; Marrino, C.L.; Marques, M.V.; Martins, J
A; Authors: Martins, E.M.F.; Matsukuma, A.Y.; Menck, C.F.M.; Miracca, E.C.; Miyaki, C.Y.
F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, D.
F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, D.
Rodriques, V.; Rosa, A.J. de M.; de Rosa Jr., V.E.; de Sa, R.G.; Santelli, R.V.; Sawasa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;anonymous, The Xylella fastidiosa Consortium of the Organization for Nucleotide Nature 406, 151-157, 2000
A;Title: The genome sequence of the plant pathogen Xylella fastidiosa.
A;Reference number: A82515; MUID:20365717; PMID:10910347
A;Note: for a complete list of authors see reference number A59328 below A;Accession: E82662
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A;Residues: 1-1784 <PYL>
A;Cross-references: UNIPROT:Q94606; GB:AE001274; NID:g3264850; PIDN:AAC24667.1;
A:Experimental source: strain MHOM/IL/81/Friedlin
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A;Title: Leishmania major Friedlin chromosome 1 has an unu
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C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999
C;Accession: H81461; T02844
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A;Cross-references: UNIPROT:Q19194; EMBL:U52002; PIDN:AAB37728.1; GSPDB:GN00022; CESP:FGA;Experimental source: strain Bristol N2; clone F08B4
C;Genetics:
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C; Accession:
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C;Species: Leishmania major
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A; Introns: 25/3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BC transporter ATP-binding protein XF1602 [imported] - Xylella fastidiosa (strain;Species: Xylella fastidiosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Molecule type: DNA
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Best Local Similarity
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25.4%;
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Pred. No. 30;
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A37845
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gamma-interferon-inducible protein IP-30 precursor - human C;Species: Homo sapiens (man) C;Date: 03-Mar-1993 #sequence_revision 03-Mar-1993 #text_change C;Accession: A43708
                                                                                                                                                                                            R;Luster, A.D.; Weinshank, R.L.; Feinman, R.; Ravetch, J. Biol. Chem. 263, 12036-12043, 1988
A;Title: Molecular and biochemical characterization of A;Reference number: A43708; MUID:88298888; PMID:3136170
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                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-303 < LUS>
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C;Accession: T26050
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                                                                                                          A;Cross-references: UNIPROT:P13284; EMBL:J03909; NID:g186264; PIDN:AAA36105.1; PID:g3070
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                                                                                                                                                         Status: preliminary
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                                       Matches
                                                                        Query Match
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Pred. No.
                                                      Pred.
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Pred. No.
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7; Mismatches
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                                       Mismatches
                                                        53.5;
No. 2
                                                                                                                                                                                                                                                R.; Ravetch, J.V.
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(EC

3.1.1.1)

precursor -

Streptomyces

scabies

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C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 C;Accession: AE2414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Streptomyces coelicolor
G;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999
C;Accession: T35335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Oliver, K.; Harris, D.; James, K.D.; Parkhill, J.; Barrell, submitted to the EMBL Data Library, May 1999
A;Reference number: Z21575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:P22266; GB:M57297; NID:g153254; PIDN:AAA26743.1; PID:g153255 R;Schottel, J.L.; Hale, V.; Babcock, M.J. Gene 115, 27-31, 1992 A;Fitle: Regulation and secretion of an extracellular esterase from Streptomyces scabies A;Reference number: PC1103; MUID:92307438; PMID:1612447 A;Accession: PC1103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       J. Bacteriol. 172, 7020-7026, 1990
A;Title: Cloning, sequencing, and regulation of expression A;Reference number: A37845; MUID:91072254; PMID:2254271 A;Accession: A37845
A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium And
                       DNA Res.
                                                                    R;Kaneko,
                                                                                                                                                                                                                                                                                                                                           S
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A;Residues: 1-54 <SCH>
A;Cross-references: GB:M57297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypothetical protein SC5H1.19c - Streptomyces coelicolor
                          Nakazaki, N.; Shimpo, S.
NA Res. 8, 205-213, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;Keywords: carboxylic ester hydrolase;1-39/Domain: signal sequence #status predicted <SIG>;40-345/Product: carboxylesterase #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Residues: 1-135 \ULL (198757; Cross-references: UNIPROT: Q9X7S7; Cross-references: Strain A3(2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   :Comment: This protein
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Accession: A37845; PC1103
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                                          T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada
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Pred. No. 24;
3; Mismatches
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Pred. No.
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                                                                                                          Anabaena sp. strain PCC 7120
#text_change 09-Jul-2004
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H
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                                          S.; Watanabe, A.; M.; Yasuda, M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B.G.; Rajandream,
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R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, .; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R. J. Bacteriol. 183, 4823-4838, 2001
                                                                                                                                                                                                                            dTDP-D-glucose 4,6-dehydratase [imported] - Clostridium acetobutylicum
C;Species: Clostridium acetobutylicum
                     A,Title: Genome Sequence and Comparative Analysis of the A,Reference number: A96900; MUID:21359325; PMID:21359325 A;Accession: B97187 A;Status: preliminary
                                                                                                                                                                               C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004 C;Accession: E97187
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A;Cross-references: UNIPROT:P96452; GB:AL591985; PIDN:CAC49304.1; PID:g15140790; GSPDB:GN
A;Experimental source: strain 1021, megaplasmid pSymB
R;Galibert, F.; Finan, T.M.; Long, S.R.; Puhler, A.; Abola, P.; Ampe, F.; Barloy-Hubler,
pela, D.; Chain, P.; Cowie, A.; Davis, R.W.; Dreano, S.; Federspiel, N.A.; Fisher, R.F.;
L.; Hyman, R.W.; Jones, T.
Science 293, 668-672, 2001
A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure,
hebault, P.; Vandenbol, M.; Vorholter, F.J.; Weidner, S.; Wells, D.H.; Wong, K.; Yeh, K.(
A;Title: The composite genome of the legume symbiont Sinorhizobium meliloti.
A;Reference number: A96039; MUID:21368234; PMID:11474104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Finan, T.M.; Weidner, S.; Wong, K.; Buhrmester, J.; Chain, P.; Vorholter, F.J.; He Proc. Natl. Acad. Sci. U.S.A. 98, 9889-9994, 2001
A;Title: The complete sequence of the 1,683-kb pSymB megaplasmid from the N2-fixing A;Reference number: A95842; MUID:21396508; PMID:11481431
A;Accession: H95954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probable Sir2-like transcription silencer protein [imported] - Sinorhizobiu
C;Species: Sinorhizobium meliloti
C;Date: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Residues: 1-192 <KUR>
A;Cross-references: UNIPROT:Q8YMR4; GB:BA000019; PIDN:BAB76568.1; PID:g17134006; GSPDB:GN
A;Experimental source: strain PCC 7120
                                                                                                                                                                                                          A;Genome: plasmid C;Superfamily: uncharacterized conserved protein with Sir2 domain
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8; Conserv
EATVLFIGYSMSDLNIRLLLHRIWSIWRRSGHEKDRP--PSYVFMY
                                                 QAQLLRVGCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
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                                                                                                      10; Mismatches
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K.S.; Zeng,

Q.; Gibson,

Solvent-Producing Bacterium Clos

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Result No.
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Maximum Match 100%
Listing first 45 summaries
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

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5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

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GenCore version 5.1.6 (c) 1993 - 2005 Compugen Ltd.
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US-08-081-863-3
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US-09-0813-345C-15
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US-09-280-501-8
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US-09-280-910-2813-34
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US-09-280-910-2813-34
US-09-280-910-2813-3139
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9374, Ap
28510, A
222132, A
32339, A
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US-09-280-501-6
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	12, Appl	4, Appli	2, Appli	31990, A	11202, A	20281, A	8072, Ap	2, Appli	32681, A	46206, A	<ol><li>Appli</li></ol>	2, Appli	2, Appli	68, Appl	68, Appl	31, Appl	14178, A	יקע, ישעי

## ALIGNMENTS

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Sequence 11, Application US/09280501

Patent No. 6440421

GENERAL INFORMATION:
APPLICANT: Cooper, Garth James Smith
APPLICANT: Reid, Ian Reginald
APPLICANT: Cornish, Jilian
TITLE OF INVENTION: TREATMENT OF BONE DISORDERS WITH
TITLE OF INVENTION: ADRENOMEDULLIN OR ADRENOMEDULLIN ACTURED APPLICATION NUMBER: US/09/280,501
CURRENT APPLICATION NUMBER: US/09/280,501
CURRENT APPLICATION NUMBER: 08/634,562
PRIOR APPLICATION NUMBER: 08/634,562
PRIOR APPLICATION NUMBER: 1990-04-18
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Best Local Similarity
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NUMBER OF SEQ ID NOS: 17
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 11
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SEQ ID NO 6
LENGTH: 38
TYPE: PRT
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CURRENT FILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: 08/634,562
PRIOR FILING DATE: 1996-04-18
NUMBER OF SEQ ID NOS: 17
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APPLICANT: Reid, Ian Reginald
APPLICANT: Cornish, Jillian
TITLE OF INVENTION: TREATMENT OF BONE DISORDERS WITH
TITLE OF INVENTION: ADRENOMEDULLIN OR ADRENOMEDULLIN
FILE REFERENCE: 08987-005001
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Pred. No. 0.0098;
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RESULT 4
US-09-813-345C-14
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US-09-280-501-11
                          GENERAL INFORMATION:
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APPLICANT:
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                                                                                                                                                                                                                                                                      TOPOLOGY: 1i
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LENGTH: 52 amino acids
TYPE: amino acid
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ADDRESSEE: Mueting, Raasch & Gebhardt, P.A.
STREET: 119 No. 6268474th Fourth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: McCormack, Myra H
REGISTRATION NUMBER: 36,602
REFERENCE/DOCKET NUMBER: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                              Local Similarity 38.
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TTLE OF INVENTION: SUPERFAMILY AND METHODS OF USE
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CREIGHTON UNIVERSITY SMITH, Derek D.
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Abel, Peter W.
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       Best Local Similarity Matches 15; Conserv
                                                                          Query Match
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (212) 596-90
TELEPAX: (212) 596-9090
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 23
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CURRENT APPLICATION NUMBER: US/09/813,345C
CURRENT FILING DATE: 2001-03-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 09/070,504
PRIOR FILING DATE: 1998-04-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: PEPTIDE ANTAGONISTS OF CGRP-RECEPTOR SUPERFAMILY AND METHODS OF TITLE OF INVENTION: USE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: HALEY Jr., James F.
REGISTRATION NUMBER: 27,794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: SHITELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                         MOLECULE TYPE:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U
ZIP: 10020
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                                                                                                                                                                                                                                                    : 185 amino acids
amino acid
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ABEL, Peter W
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MATSUO, Hisayuki
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Score 72.5; DB Pred. No. 0.056; 6; Mismatches
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                                                                      DB 1;
                                                                      Length 185;
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US-08-801-863-1
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                                                                                                                                                                                                                                                                                                                              Matches
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/801,863
FILING DATE: CONCURRENTLY HEREWITH
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: HALEY Jr., James F.
REFERENCE/DOCKET NUMBER: S17,794
REFERENCE/DOCKET NUMBER: SHGN-5 DIV3
TELECOMMUNICATION INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 185 amino acids
                                                                                                                                                 GENERAL INFORMATION:
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APPLICANT: KITAMU
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TOPOLOGY: lin-
OLECULE
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                TITLE OF INVENTION: ADRENOMEDULLIN NUMBER OF SEQUENCES: 10 CORRESPONDENCE ADDRESS:
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STREET:
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                ADDRESSEE:
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Similarity 38.5%;
15; Conservative
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 1251 Avenue of the Americas
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1251 Avenue of the Americas
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(212) 596-000
                                                                             KANGAWA, Kenji
MATSUO, Hisayuki
ETO, Tanenao
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MATSUO, Hisayuki
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                    c/o FISH & NEAVE
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                                                                                                                                                                                                                                                                                                                            6; Mismatches
                                                                                                                                                                                                                                                                                                                         17; Indels
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RESULT 8
US-09-004-713-1
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                                    TELEFAX: (212) 596-9090 INFORMATION FOR SEQ ID NO:
                                                                                   APPLICATION NUMBER: US/09/004
FILING DATE: JANUARY 7, 1998
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: HALEY Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: SHGN
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: KITAMU
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 185 amino acids
                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
SEQUENCE CHARACTERISTICS:
LENGTH: 185 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: HALEY Jr., James F.
REGISTRATION NUMBER: 27,794
REGISTRATION NUMBER: 27,794
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TELECOMMUNICATION INFORMATION:
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TITLE OF INVENTION:
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TOPOLOGY: linear
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                                                                      TELEPHONE:
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1251 Avenue of the Americas
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MATSUO, Hisayuki
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38.5%;
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                                                                                                            SHGN-5 DIV2 CON
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; TOPOLOGY: linear ; MOLECULE TYPE: protein US-09-004-713-1

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US-08-801-863-3
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                                                                                                RESULT 10
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Best Local Similarity 38.5%;
Matches 15; Conservative
                                                   equence 3, Application US/08801863
                                                                                                                                                                                                                                                                                          TELEFAX: (212) 596-9090 INFORMATION FOR SEQ ID NO: 3:
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                                                                                                                                                                                                                                 MOLECULE TYPE: protein
                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 188 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/08233389C
Patent No. 5639855
                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 27,794
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 38.5%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                      NAME: HALEY Jr., Jam
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: LECTY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 26-APP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: KITAMURA, Kazuo
APPLICANT: KANGAWA, Kenji
APPLICANT: MATSUO, Hisayuki
                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                            109 ĠĊŖŖĠŦĊŦŸŎĸĹĄĦŊŢŶŖŦ-DĸDĸDĠŸĀŔŖĸĸĬŚŔŊĠŶ 146
                                            INFORMATION:
                                                                                                                                 9 GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDDSSPHSY 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TLE OF INVENTION: ADRENOMEDULLIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MBER OF SEQUENCES:
KITAMURA, Kazuo
KANGAWA, Kenji
MATSUO, Hisayuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                109 ĠĆRFĠŤĊTŸŎKĹAHQIYÓFT-DKDKDNVÁFRSKIŠPQGÝ 146
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: C/O FISH & NEAVE
1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                       26-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tanenao
                                                                                                                                                                                                                                                                                                                                                                  James F
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                                                                                                                                                                                                                                                                                                                                                                                                          US/08/233,389C
                                                                                                                                                         Score 70.5; DB 1; Length 188; Pred. No. 0.11; Length 188; S; Mismatches 18; Indels
                                                                                                                                                                                                                                                                                                                                       SHGN-5
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                       STREET: 120.
CITY: New York
CTATE: NY
                                                                                                                                                                                                                                                                                                         08-486-596A-3
                                                                                                     STALL.
COUNTRY: US.
10020
                                                                                                                                                                       ADDRESSEE:
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ATTORNEY AGENT INFORMATION:
NAME: HALEY JI., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: SHGN-5 DIVI
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GENERAL INFORMATION:
                                                                                                                                                                 COMPUTER: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION DATA:
APPLICATION NUMBER: US/08/486,596A
CLASSIFICATION: 530
APPRIVATION: 530
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 188 amino acids
TYPE: amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: KITAMURA, Kazuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
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REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: SHI
TELECOMMUNICATION INFORMATION.
(212) 596-9000
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OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
PATENTA PAPLICATION DATA:
FILING DATA:
CLASSIPTOSE:
CONCIDEDENT.
CLASSIPTOSE:
CONCIDEDENT.
CON
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ADDRESSEE: C/O FISH & NEAVE
STREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: LEGISTREET: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          109 ĠĆRFĠŤĊŢVQKĹAHQIYQFT-DKDKDGVÁĖRSKIŚĖQGY 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3, Application US/08486596A
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15; Conservative
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KANGAWA, Kenj
MATSUO, Hisayuki
ETO, Tanenao
ADRENOMF
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51 Avenue of the Americas
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38.5%;
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Pred. No. 0.11;
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Gaps

TELECOMMUNICATION INFORMATION:

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RESULT 13
US-09-070-504-15
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US-09-004-713-3
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                                                                                                                                                                                               ; MOLECULE TYPE: protein US-09-004-713-3
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                                                                                                                                Matches
                                                                                                                                                Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                 FILING DATE: JANUARY 7, 1998
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: HALEY Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: SHGN-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                TELEFAX: (212) 596-9090 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212) 596-9090 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: ETO, Tane
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                 TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: New York
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                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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                                                                                                                                                                                                                                                                  ENGTH:
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                                                               109 GCRFGTCTVQKLAHQIYQFT-DKDKDGVAPRSKISPQGY 146
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al Similarity 38.5%;
15; Conservative
                                                                                             9 GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
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                                                                                                                                                h 27.8%;
Similarity 38.5%;
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1251 Avenue of the Americas
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                                                                                                                                  Conservative
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MATSUO, Hisayuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KITAMURA,
                                                                                                                                                                                                                                                                                                                                     (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tanenao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADRENOMEDULLIN
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                                                                                                                                                Score 70.5; DB 2; Length 188; Pred. No. 0.11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 70.5; DB Pred. No. 0.11;
                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18; Indels
                                                                                                                              18; Indels
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US-09-813-345C-15
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                                        SOFTWARE: PatentIn version 3.2
SEQ ID NO 15
LENGTH: 50
                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: CREIGHTON UNIVERSITY
APPLICANT: SMITH, Derek D.
APPLICANT: SAHA, Shankar
APPLICANT: ABEL, Peter W.
                                                                                                                                                                                                                                                                                                                                                                           Sequence 15, Application US/09813345C Patent No. 6756205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                     FILE REFERENCE: 180.00020102
CURRENT APPLICATION NUMBER: US/09/813,345C
CURRENT FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: 09/070,504
PRIOR FILING DATE: 1998-04-30
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                             TITLE OF INVENTION: PEPTIDE ANTAGONISTS OF CGRP-RECEPTOR SUPERFAMILY AND METHODS OF TITLE OF INVENTION: USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 612/305-1228
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/070,504
FILING DATE: 30-APR-1998
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 36,602
REFERENCE/DOCKET NUMBER: 180.00020101
TELECOMMUNICATION INFORMATION:
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NAME: McCormack, Myra H
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Saha, Shankar APPLICANT: Abel, Peter W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: PEPTIDE ANTAGONI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 27.4%;
Local Similarity 35.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 119 No. 6:
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9 GCVLGTCQVQNLSHRLWQLMGPAGRQDSAPVDPSSPHSY 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           55401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCRFGTCTMQKLAHQIYQFT-DKDKDGMAPRNKISPQGY 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              50 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mueting, Raasch & Gebhardt, P.A.
19 No. 6268474th Fourth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              612/305-1220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Derek D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PEPTIDE ANTAGONISTS OF CGRP-RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 69.5; DB 3; Length 50; Pred. No. 0.033;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ۲.
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OTHER IMPORMATION: Artificially Synthesized Peptide
US-09-813-345C-15

Query Match
Best Local Similarity 35.94; Pred, No. 0.03;
Matches 14; Conservative 7; Mismatches 17; Indels 1; Gaps 1;

Qy 9 GCVITCOVONISHRUMOLMGPAGRODSAPVDSSPHS 47
Db 13 GCREGTCTMOKLAHGITOFFT-DKDKDGMAPRNKISPQGY 50

RESULT 15
US-09-880-501-8
US-09-880-501-8
US-09-880-501-8
US-09-880-501-8
US-09-880-501-8
INFORMATION: APPLICANTON GARTH James Smith
APPLICANT: Cooper, Garth James Smith
ITTLE OF INVENTION: TREATMORT OF BOME DISORDERS WITH
ITTLE OF INVENTION: ADERENOMEDULLIN OR ADREMOMEDULLIN AGONISTS
FILE REFERENCE: 08997-050.01
CURRENT APPLICATION NUMBER: US/09/280,501
PRIOR APPLICATION NUMBER: US/09/280,501
PRIOR APPLICATION NUMBER: US/09/280,501
FRIDE APPLICANTON NUMBER: US/09/280,501
FRICK APPLICATION NUMBER: US/09/280,501
FRICK APPLICANTON NUM
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